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(54) **PHARMACEUTICAL COMPOSITION FOR TREATMENT AND/OR PROPHYLAXIS OF CANCER**

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(56) **References Cited**

U.S. PATENT DOCUMENTS

5,698,396 A 12/1997 Pfreundschuh  
6,335,170 B1 1/2002 Orntoft  
6,444,425 B1 9/2002 Reed et al.  
7,449,184 B2 11/2008 Allison et al.

7,485,302 B2 2/2009 Adams et al.  
7,745,391 B2 6/2010 Mintz et al.  
8,211,634 B2 7/2012 Depinho et al.  
8,709,418 B2 4/2014 Okano et al.  
8,828,398 B2 9/2014 Kobayashi et al.  
8,911,740 B2 12/2014 Saito et al.  
2002/0006404 A1 1/2002 Hanna et al.  
2003/0118599 A1 6/2003 Algate et al.  
2003/0190640 A1 10/2003 Faris et al.  
2004/0029114 A1 2/2004 Mack et al.  
2004/0236091 A1 11/2004 Chicz et al.  
2004/0258678 A1 12/2004 Bodary et al.  
2005/0003390 A1 1/2005 Axenovich et al.  
2005/0032113 A1 2/2005 Tanaka et al.  
2005/0244413 A1 11/2005 Adolf et al.  
2006/0019256 A1 1/2006 Clarke et al.  
2006/0069054 A1 3/2006 Houghton et al.  
2006/0275305 A1 12/2006 Bryant  
2007/0048301 A1 3/2007 Bodary-Winter et al.  
2007/0154931 A1 7/2007 Radich et al.  
2007/0264253 A1 11/2007 Liu et al.  
2008/0075722 A1 3/2008 DePinho et al.  
2008/0107668 A1 5/2008 Philip et al.  
2010/0068724 A1 3/2010 Fung et al.

(Continued)

FOREIGN PATENT DOCUMENTS

CN 1705676 A 12/2005  
CN 101120252 A 2/2008

(Continued)

OTHER PUBLICATIONS

Bodey et al., "MAGE-1, a Cancer/Testis-Antigen, Expression in Childhood Astrocytomas as an Indicator of Tumor Progression," in vivo (2002) vol. 16, pp. 583-588.

(Continued)

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(57)

**ABSTRACT**

An object of the present invention is to prepare an antibody that targets CAPRIN-1 specifically expressed on the surface of cancer cells and is superior in antitumor activity to conventional antibodies and to provide use of the antibody as a therapeutic and/or preventive agent for cancer. The present invention provides use of an antibody targeting an identified cancer antigenic protein specifically expressed on the surface of cancer cells as a therapeutic and/or preventive agent for cancer, specifically, a pharmaceutical composition for treatment and/or prevention of cancer, comprising as an active ingredient an antibody or a fragment thereof which has immunological reactivity with a CAPRIN-1 protein, the antibody or the fragment thereof comprising a heavy chain variable region comprising amino acid sequences represented by SEQ ID NOs: 5, 6, and 7 and a light chain variable region comprising amino acid sequences represented by SEQ ID NOs: 9, 10, and 11.

**15 Claims, No Drawings**

(56)

**References Cited****U.S. PATENT DOCUMENTS**

2011/0123492	A1	5/2011	Okano et al.
2011/0136121	A1	6/2011	Okano et al.
2011/0189700	A1	8/2011	Moses et al.
2011/0256144	A1	10/2011	Okano et al.
2012/0171699	A1	7/2012	Goodman et al.
2012/0214975	A1	8/2012	Sandig et al.
2012/0294860	A1	11/2012	Ido et al.
2012/0301471	A1	11/2012	Kobayashi et al.
2012/0301476	A1	11/2012	Okano et al.
2012/0321641	A1	12/2012	Okano et al.
2013/0045210	A1	2/2013	Kobayashi et al.
2013/0071398	A1	3/2013	Saito et al.

**FOREIGN PATENT DOCUMENTS**

CN	101189516	A	5/2008
CN	102170907	A	8/2011
EP	2 325 648	A1	5/2011
EP	2322221	A1	5/2011
EP	2 532 367	A1	12/2012
EP	2 532 743	A1	12/2012
EP	2 832 365	A1	2/2015
EP	2 832 366	A1	2/2015
JP	2002-540790	A	12/2002
JP	2003-528587	A	9/2003
JP	2006-316040	A	11/2006
JP	2013-502205	A	1/2013
JP	2013-505028	A	2/2013
RU	2234942	C2	2/2003
RU	2306952	C2	9/2007
RU	2006137060	A	4/2008
WO	WO 00/04149	A2	1/2000
WO	WO 00/60077	A2	10/2000
WO	WO 01/32910	A2	5/2001
WO	WO 01/72295	A2	10/2001
WO	WO 02/078524	A2	10/2002
WO	WO 02/083070	A2	10/2002
WO	WO 02/092001	A2	11/2002
WO	WO 2004/076682	A2	9/2004
WO	WO 2004/097051	A2	11/2004
WO	WO 2005/007830	A2	1/2005
WO	WO 2005/100998	A2	10/2005
WO	WO 2005/116076	A2	12/2005
WO	WO 2006/002378	A2	1/2006
WO	WO 2007/150077	A2	12/2007
WO	WO 2008/031041	A2	3/2008
WO	WO 2008/059252	A2	5/2008
WO	WO 2008/073162	A2	6/2008
WO	WO 2008/088583	A2	7/2008
WO	WO 2009/113742	A1	9/2009
WO	WO 2010/016525	A1	2/2010
WO	WO 2010/016526	A1	2/2010
WO	WO 2010/016527	A1	2/2010
WO	WO 2011/096517	A1	8/2011
WO	WO 2011/096528	A1	8/2011
WO	WO 2011/096533	A1	8/2011
WO	WO 2011/096534	A1	8/2011
WO	WO 2011/096535	A1	8/2011
WO	WO 2013/018885	A1	2/2013
WO	WO 2013/018886	A1	2/2013
WO	WO 2013/018894	A1	2/2013
WO	WO 2013/147169	A1	10/2013
WO	WO 2013/147176	A1	10/2013

**OTHER PUBLICATIONS**

Comtesse et al., "Probing the human natural autoantibody repertoire using an immunoscreening approach," *Clin. Exp. Immunol.* (2000), vol. 121, pp. 430-436.

International Search Report issued Nov. 18, 2014, in PCT International Application No. PCT/JP2014/071094.

Jager et al., "Identification of a Tissue-specific Putative Transcription Factor in Breast Tissue by Serological Screening of a Breast Cancer Library," *Cancer Research* (Mar. 1, 2001), vol. 61, pp. 2055-2061.

Jungbluth et al., "Immunohistochemical Analysis of NY-ESO-1 Antigen Expression in Normal and Malignant Human Tissues," *Int. J. Cancer* (2001), vol. 92, pp. 856-860.

Kohler et al., "Tumor antigen analysis in neuroblastoma by serological interrogation of bioinformatic data," *Cancer Science* (Nov. 2010), vol. 101, No. 11, pp. 2316-2324.

Nakamura et al., "Gene Expression Profile of Metastatic Human Pancreatic Cancer Cells Depends on the Organ Microenvironment," *Cancer Research* (Jan. 1, 2007), vol. 67, No. 1, pp. 139-148.

Non-Final Office Action issued Nov. 6, 2014, in U.S. Appl. No. 13/576,950.

Pegram et al., "Rational Combinations of Trastuzumab with Chemotherapeutic Drugs Used in the Treatment of Breast Cancer," *Journal of the National Cancer Institute* (May 19, 2004), vol. 96, No. 10, pp. 739-749.

Punt et al., "Edrecolomab alone or in combination with fluorouracil and folinic acid in the adjuvant treatment of stage III colon cancer: a randomised study," *Lancet* (Aug. 31, 2002), vol. 360, No. 9334, pp. 671-677.

Buchsbaum et al., "Treatment of Pancreatic Cancer Xenografts with Erbitux (IMC-C225) Anti-EGFR Antibody, Gemcitabine, and Radiation," *Int. J. Radiation Oncology Biol. Phys.* (2002), vol. 54, No. 4, pp. 1180-1193.

Chames et al., "Therapeutic Antibodies for the Treatment of Pancreatic Cancer," *The Scientific World Journal* (Jan. 1, 2010), vol. 10, pp. 1107-1120.

Eccleston et al., "Pancreatic Tumor Marker Anti-Mucin Antibody CAM 17.1 Reacts with a Sialyl Blood Group Antigen, Probably I, Which is Expressed throughout the Human Gastrointestinal Tract," *Digestion* (1998), vol. 59, pp. 665-670.

Esteva et al., "Chemotherapy of Metastatic Breast Cancer: What to Expect in 2001 and Beyond," *The Oncologist* (2001), vol. 6, pp. 133-146.

Extended European Search Report issued Feb. 2, 2015, in European Patent Application No. 12819473.5.

Extended European Search Report issued Jan. 29, 2015, in European Patent Application No. 12819899.1.

Houghton, P. J. and J. A. Houghton, "Evaluation of Single-Agent Therapy in Human Colorectal Tumour Xenografts," *Br. J. Cancer* (1978), vol. 37, pp. 833-840.

GenBank Accession No. AAU93399, Sep. 22, 2005.

GenBank Accession No. BAF96513, Jan. 5, 2008.

GenBank Accession No. NM\_001031365, Sep. 25, 2007.

GenBank Accession No. NM\_001076062, Feb. 9, 2008.

GenBank Accession No. NM\_001111289, Feb. 11, 2008.

GenBank Accession No. NM\_001111290, Feb. 11, 2008.

GenBank Accession No. NM\_001111291, Feb. 10, 2008.

GenBank Accession No. NM\_001111292, Feb. 11, 2008.

GenBank Accession No. NM\_016739, Feb. 10, 2008.

GenBank Accession No. NM\_05898, Feb. 11, 2008.

GenBank Accession No. NM\_203364, Feb. 10, 2008.

GenBank Accession No. Q14444, Jun. 10, 2008.

GenBank Accession No. Q1LZB6, Jun. 10, 2008.

GenBank Accession No. XM\_853016, Aug. 30, 2005.

Patent Examination Report No. 1 issued Oct. 14, 2014, in Australian Patent Application No. 2009278387.

Akiyoshi, "Cancer Vaccine Therapy Using Peptides Derived from Tumor-Rejection Antigens," *Jpn J Cancer Chemother.*, vol. 24, No. 5, Mar. 1997, pp. 511-519, with English Abstract (p. 519).

Balmana et al., "BRCA in breast cancer: ESMO Clinical Recommendations," *Annals of Oncology*, vol. 20, Supplemental 4, 2009, pp. iv19-iv20.

Bodey et al., "Failure of Cancer Vaccines: The Significant Limitations of this Approach to Immunotherapy," *Anticancer Research*, vol. 20, 2000, pp. 2665-2676.

Brand et al., "Prospect for Anti-HER2 Receptor Therapy in Breast Cancer," *Anticancer Research*, vol. 26, 2006, pp. 463-470.

Brass et al., "Translation initiation factor eIF-4gamma is encoded by an amplified gene and induces an immune response in squamous cell lung carcinoma," *Human Molecular Genetics*, vol. 6, No. 1, 1997, pp. 33-39.

(56)

## References Cited

## OTHER PUBLICATIONS

- Chamberlain et al., "Innovations and strategies for the development of anticancer vaccines," *Expert Opinion on Pharmacotherapy*, vol. 1, No. 4, 2000, pp. 603-614.
- Ellis, et al., "Identification and Characterization of a Novel Protein (p137) Which Transcytoses Bidirectionally in Caco-2 Cells", *The Journal of Biological Chemistry*, Sep. 1, 1995, vol. 270, No. 35, pp. 20717-20723.
- European Search Report, dated Aug. 13, 2013, for European Application No. 11739882.6.
- European Search Report, dated Aug. 26, 2011, for European Application No. No. 09805010.7.
- European Search Report, dated Jan. 30, 2013, for European Application 09805009.9.
- European Search Report, dated Nov. 6, 2013, for European Application No. 11739876.8.
- Evans et al., "Vaccine therapy for cancer—fact or fiction?", *Q J Med*, vol. 92, 1999, pp. 299-307.
- GeneCards, "Cell Cycle Associated Protein 1—Biological research products for CAPRIN 1," updated Mar. 19, 2013, 10 pages.
- Grill et al., "Activation/Division of Lymphocytes Results in Increased Levels of Cytoplasmic Activation/Proliferation-Associated Protein-1: Prototype of a New Family of Proteins," *The Journal of Immunology*, vol. 172, 2004, pp. 2389-2400.
- Güre et al., "Human Lung Cancer Antigens Recognized by Autologous Antibodies: Definition of a Novel cDNA Derived from the Tumor Suppressor Gene Locus on Chromosome 3p21.3," *Cancer Research*, vol. 58, Mar. 1, 1998, pp. 1034-1041.
- Güre et al., "SSX: A Multigene Family with Several Members Transcribed in Normal Testis and Human Cancer," *Int. J. Cancer*, vol. 72, 1997, pp. 965-971.
- Harlow et al., "Antibodies a Laboratory Manual", Cold Spring Harbor Laboratory, Chapter 3, 1988, pp. 23-34.
- Hugo Gene Nomenclature Committee, Gene Symbol Report, CAPRIN1, Approved Name: Cell Cycle Associated Protein 1, HGNC ID: HGNC:6743, Nov. 3, 2012, 2 pages.
- International Search Report and Written Opinion of the International Searching Authority, dated Mar. 1, 2011, for International Application No. PCT/JP2011/052413.
- International Search Report, dated Mar. 15, 2011, for International Application No. PCT/JP2011/052384.
- International Search Report, dated Mar. 8, 2011, for International Application No. PCT/JP2011/052403.
- International Search Report, dated Mar. 8, 2011, for International Application No. PCT/JP2011/052414.
- International Search Report, dated Oct. 6, 2009, for International Application No. PCT/JP2009/063882.
- International Search Report, dated Sep. 8, 2009, for International Application No. PCT/JP2009/063883.
- Itoh et al., "HUB1 is an autoantigen frequently eliciting humoral immune response in patients with adult T cell leukemia," *Int. J. Oncol.*, vol. 14, No. 4, Apr. 1999, pp. 703-708 (Abstract only provided).
- Jang et al., "Antihypertensive Angiotensin I-Converting Enzyme Inhibitory Activity and Antioxidant Activity of Vitis hybrid-Vitis coignetiae Red Wine Made with Saccharomyces cerevisiae," *Mycobiology*, vol. 39, No. 2, 2011, pp. 137-139.
- Kaddar et al., "Two new miR-16 targets: caprin-1 and HMGA1, proteins implicated in cell proliferation," *Biology of the Cell*, vol. 101, No. 9, 2009, pp. 511-524.
- Kajiji et al., "Six Monoclonal Antibodies to Human Pancreatic Cancer Antigens," *Cancer Research*, vol. 47, Mar. 1, 1987, pp. 1367-1376.
- Karazum et al., "Caprin 1 is Frequently Overexpressed in Human Lymphomas," *American Society of Human Genetics, Cancer Genetics*, Program No. 1190W, Oct. 12, 2011, One page (Abstract only).
- Kataja et al., "Primary breast cancer: ESMO Clinical Recommendations for diagnosis, treatment and follow-up," *Annals of Oncology*, vol. 20, Supplement 4, 2009, pp. iv10-iv14.
- Katsafanas et al., "Colocalization of Transcription and Translation within Cytoplasmic Poxvirus Factories Coordinates Viral Expression and Subjugates Host Functions," *Cell Host & Microbe*, vol. 2, Oct. 2007, pp. 221-228.
- Katsafanas et al., "Vaccinia Virus Intermediate Stage Transcription is Complemented by Ras-GTPase-activating Protein SH3 Domain-binding Protein (G3BP) . . .," *Jour. of Biol. Chem.*, vol. 279, No. 50, Dec. 10, 2004, pp. 52210-52217.
- Kolobova et al., "Microtubule-dependent association of AKAP350A and CCAR1 with RNA stress granules," *Experimental Cell Research*, vol. 315, 2009 (Available online Dec. 3, 2008), pp. 542-555.
- Lu et al., "Identification of an immunological signature of tumor rejection in the neu transgenic mouse," 2007 AACR Annual Meeting, Apr. 14-18, 2007 (Presentation conducted on Apr. 17, 2007), One page (Abstract only provided).
- Lu et al. "Targeting serum antibody for cancer diagnosis: a focus on colorectal cancer," *Expert Opin. Ther. Targets*, vol. 11, No. 2, 2007, pp. 235-244.
- Müller-Pillasch et al., "Identification of a new tumour-associated antigen TM4SF5 and its expression in human cancer," *Gene*, vol. 208, 1998, pp. 25-30.
- Munodzana et al., "Conformational Dependence of Anaplasma marginale Major Surface Protein 5 Surface-Exposed B-Cell Epitopes", *Infection and Immunity*, vol. 66, No. 6, Jun. 1998, pp. 2619-2624.
- NCBI Reference Sequence, caprin-1 [Bos taurus], 2009, Accession No. NP\_001069530, XP\_615677, 1 page.
- NCBI Reference Sequence, caprin-1 [Gallus gallus], 2005, Accession No. NP\_001026536, XP\_423820, 1 page.
- NCBI Reference Sequence, caprin-1 isoform 1 [Homo sapiens], 1995, Accession No. NP\_005889, 3 pages.
- NCBI Reference Sequence, caprin-1 isoform 2 [Homo sapiens], 1995, Accession No. NP\_976240, 3 pages.
- NCBI Reference Sequence, caprin-1 isoform a [Mus musculus], 1996, Accession No. NP\_058019, 3 pages.
- NCBI Reference Sequence, caprin-1 isoform b [Mus musculus], 1996, Accession No. NP\_001104760, 3 pages.
- NCBI Reference Sequence, caprin-1 isoform c [Mus musculus], 1996, Accession No. NP\_001104761, 4 pages.
- NCBI Reference Sequence, Predicted: caprin-1 [Equus caballus], 2008, Accession No. XP\_001492799, 1 page.
- NCBI Reference Sequence, Predicted: caprin-1 isoform 2 [Canis lupus familiaris], Dec. 2, 2011, Accession No. XP\_858109, 1 page.
- Nelson et al., "Screening for Breast Cancer: An Update for the U.S. Preventive Services Task Force," *Ann. Intern. Med.*, vol. 151, No. 10, Nov. 17, 2009, pp. 727-737.
- Okano et al., "Abstract 519: Identification of a novel target for antibody therapy of breast cancer", *Cancer Research*, vol. 72, Issue 8, Supplement 1, Apr. 15, 2012, XP-002700046, 2 pages.
- Polyak et al., "Alanine-170 and proline-172 are critical determinants for extracellular CD20 epitopes; heterogeneity in the fine specificity of CD20 monoclonal antibodies is defined by additional requirements imposed by both amino acid sequence . . .", *Blood*, vol. 99, No. 9, May 1, 2002, pp. 3256-3262.
- R & D Systems, "IHC Products & Protocol Guide," printed Jan. 9, 2014, pp. 1-112.
- Sahin et al., "Human neoplasms elicit multiple specific immune responses in the autologous host," *Proc. Natl. Acad. Sci. USA*, vol. 92, Dec. 1995, pp. 11810-11813.
- Scanlan et al., "Cancer-related Serological Recognition of Human Colon Cancer: Identification of Potential Diagnostic and Immunotherapeutic Targets," *Cancer Research*, vol. 62, Jul. 15, 2002, pp. 4041-4047.
- Scanlan et al., "Characterization of Human Colon Cancer Antigens Recognized by Autologous Antibodies," *Int. J. Cancer*, vol. 76, 1998, pp. 652-658.
- Solomon et al., "Distinct Structural Features of Caprin-1 Mediate Its Interaction with G3BP-1 and Its Induction of Phosphorylation of Eukaryotic Translation Initiation Factor 2 $\alpha$ , Entry to Cytoplasmic Stress . . .," *Molecular and Cellular Biology*, vol. 27, No. 6, Mar. 2007, XP\_002690351, pp. 2324-2342.

(56)

**References Cited**

## OTHER PUBLICATIONS

Strome et al., "A Mechanistic Perspective of Monoclonal Antibodies in Cancer Therapy Beyond Target-Related Effects," *The Oncologist*, vol. 12, 2007, pp. 1084-1095.

Türeci et al., "The SSX-2 Gene, Which Is Involved in the t(X; 18) Translocation of Synovial Sarcomas, Codes for the Human Tumor Antigen HOM-MEL-40," *Cancer Research*, vol. 56, Oct. 15, 1996, pp. 4766-4772.

United States Notice of Allowance, dated Dec. 2, 2013, for U.S. Appl. No. 13/576,955.

United States Office Action, dated Aug. 19, 2013, for U.S. Appl. No. 13/576,955.

United States Office Action, dated Aug. 26, 2013, for U.S. Appl. No. 13/576,950.

United States Office Action, dated Dec. 21, 2012, for U.S. Appl. No. 13/057,515.

United States Office Action, dated Dec. 21, 2012, for U.S. Appl. No. 13/057,709.

United States Office Action, dated Jan. 16, 2014, for U.S. Appl. No. 13/057,515.

United States Office Action, dated Jul. 1, 2013, for U.S. Appl. No. 13/057,515.

United States Office Action, dated Jul. 1, 2013, for U.S. Appl. No. 13/577,212.

United States Office Action, dated Jul. 16, 2013, for U.S. Appl. No. 13/057,709.

United States Office Action, dated Jun. 14, 2013, for U.S. Appl. No. 13/576,969.

United States Office Action, dated Mar. 13, 2013, for U.S. Appl. No. 13/576,955.

United States Office Action, dated Nov. 15, 2013, for U.S. Appl. No. 13/576,953.

United States Office Action, dated Nov. 15, 2013, for U.S. Appl. No. 13/577,028.

United States Office Action, dated Nov. 15, 2013, in U.S. Appl. No. 13/576,950.

United States Office Action, dated Nov. 2, 2012, for U.S. Appl. No. 13/057,709.

United States Office Action, dated Nov. 9, 2012, for U.S. Appl. No. 13/057,515.

United States Office Action, dated Oct. 15, 2013, for U.S. Appl. No. 13/576,969.

United States Office Action, dated Oct. 21, 2013, for U.S. Appl. No. 13/577,212.

United States Office Action, dated Sep. 19, 2013, for U.S. Appl. No. 13/577,028.

United States Office Action, dated Sep. 6, 2013, for U.S. Appl. No. 13/576,953.

Van Der Bruggen et al., "A Gene Encoding an Antigen Recognized by Cytolytic T Lymphocytes on a Human Melanoma," *Science*, vol. 254, Dec. 13, 1991, pp. 1643-1647 (Also published in *J. Immunol.*, vol. 178, 2007, pp. 2617-2621).

Wang et al., "Absence of Caprin-1 Results in Defects in Cellular Proliferation", *The Journal of Immunology*, 2005, vol. 175, pp. 4274-4282.

Yanai et al., "Dlk-1, a cell surface antigen on foetal hepatic stem/progenitor cells, is expressed in hepatocellular, colon, pancreas and breast carcinomas at a high frequency," *The Journal of Biochemistry*, vol. 148, No. 1, 2010 (Publ. online Mar. 30, 2010), pp. 85-92.

De Pascalis et al., "Grafting of "Abbreviated" Complementarity-Determining Regions Containing Specificity-Determining Residues Essential for Ligand Contact to Engineer a Less Immunogenic Humanized Monoclonal Antibody," *J. Immunol.* (2002), vol. 169, pp. 3076-3084.

Extended European Search Report issued Mar. 2, 2015, in European Patent Application No. 12819759.7.

Office Action issued Jan. 28, 2015, in Russian Patent Application No. 2012137502, with partial English translation.

Riechmann et al., "Reshaping human antibodies for therapy," *Nature* (Mar. 24, 1988), vol. 332, pp. 323-327.

Vajdos et al., "Comprehensive Functional Maps of the Antigen-binding Site of an Anti-ErbB2 Antibody Obtained with Shotgun Scanning Mutagenesis," *J. Mol. Biol.* (2002), vol. 320, pp. 415-428.

Extended European Search Report issued Mar. 18, 2015, in European Patent Application No. 12820225.6.

Extended European Search Report issued Mar. 23, 2015, in European Patent Application No. 12820596.0.

Non-Final Office Action issued Apr. 14, 2015, in U.S. Appl. No. 14/236,793.

GenBank Accession No. NM\_005898, Feb. 11, 2008.

Gong et al., "Caprin-1 is a novel microRNA-223 target for regulating the proliferation and invasion of human breast cancer cells," *Bio-medicine & Pharmacotherapy* (2013), vol. 67, pp. 629-636.

Qui et al., "Targeting a ribonucleoprotein complex containing the caprin-1 protein and the c-Myc mRNA suppresses tumor growth in mice: an identification of a novel oncotarget," *Oncotarget* (2014), vol. 6, No. 4, pp. 2148-2163.

Sabile et al., "Caprin-1, a novel Cyr61-interacting protein, promotes osteosarcoma tumor growth and lung metastasis in mice," *Biochimica et Biophysica Acta* (2013), vol. 1832, pp. 1173-1182.

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# PHARMACEUTICAL COMPOSITION FOR TREATMENT AND/OR PROPHYLAXIS OF CANCER

## TECHNICAL FIELD

The present invention relates to novel use of an antibody against CAPRIN-1 or a fragment thereof in a drug such as a therapeutic and/or preventive agent for cancer.

## BACKGROUND ART

Cancer is the leading cause of death. This disease is currently treated principally by surgical therapy in combination with radiation therapy and/or chemotherapy. In spite of recent development of novel surgical techniques or discovery of novel anticancer agents, the existing treatment of cancer has an insufficiently improved outcome, except for some cancer types. With recent advances of molecular biology or cancer immunology, antibodies that specifically react with cancer, cancer antigens that are recognized by cytotoxic T cells, genes encoding such cancer antigens, and the like have been identified, raising expectations on specific cancer therapy targeting the cancer antigens (Non Patent Literature 1).

For reducing the side effect of cancer therapy, it is desired that peptides, polypeptides, or proteins recognized as antigens of the cancer should rarely exist in normal cells and specifically exist in cancer cells. In 1991, Boon et al. (Ludwig Institute for Cancer Research, Belgium) isolated a human melanoma antigen MAGE1 recognized by CD8-positive T cells by a cDNA expression cloning method using autologous cancer cell lines and cancer-reactive T cells (Non Patent Literature 2). Then, a SEREX (serological identification of antigens by recombinant expression cloning) method has been reported, which adopts a gene expression cloning approach to identify tumor antigens recognized by antibodies produced in response to autologous cancer in vivo in a cancer patient (Non Patent Literature 3 and Patent Literature 1). According to this method, some cancer antigens that are rarely expressed in normal cells and are specifically expressed in cancer have been isolated (Non Patent Literatures 4 to 9). In addition, cell therapy using immunocytes that specifically react with cancer antigens or cancer-specific immunotherapy using vaccines or the like comprising cancer antigens is under clinical trial targeting some of the isolated cancer antigens.

In recent years, various antibody drugs for cancer treatment targeting antigenic proteins on cancer cells have emerged in the world. These drugs have received attention because of their certain efficacy as cancer-specific therapeutic agents. A large majority of antigenic proteins targeted by the drugs, however, are also expressed in normal cells. As a result of administering the antibodies, normal cells expressing the antigens as well as cancer cells are damaged, disadvantageously resulting in side effect. Thus, if cancer antigens specifically expressed on the surface of cancer cells can be identified and antibodies targeting the antigens can be used as drugs, these antibody drugs can be expected to achieve treatment with less side effect.

Cytoplasmic-activation- and proliferation-associated protein 1 (CAPRIN-1) has been known as an intracellular protein that is expressed upon activation or cell division of resting normal cells and forms cytoplasmic stress granules with intracellular RNAs to participate in the regulation of transport and translation of mRNAs. This protein has been found to be specifically expressed on the surface of cancer cells and is

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therefore under study as a target of antibody drugs for cancer treatment (Patent Literature 2).

## CITATION LIST

### Patent Literature

Patent Literature 1: U.S. Pat. No. 5,698,396

Patent Literature 2: WO2010/016526

### Non Patent Literature

Non Patent Literature 1: Tsuyoshi Akiyoshi, "Japanese Journal of Cancer and Chemotherapy", 1997, Vol. 24, p. 511-519 (Japanese Journal of Cancer and Chemotherapy Publishers Inc., Japan)

Non Patent Literature 2: Bruggen P. et al., Science, 254: 1643-1647(1991)

Non Patent Literature 3: Proc. Natl. Acad. Sci. USA, 92: 11810-11813(1995)

Non Patent Literature 4: Int. J. Cancer, 72: 965-971(1997)

Non Patent Literature 5: Cancer Res., 58: 1034-1041(1998)

Non Patent Literature 6: Int. J. Cancer, 29: 652-658(1998)

Non Patent Literature 7: Int. J. Oncol., 14: 703-708(1999)

Non Patent Literature 8: Cancer Res., 56: 4766-4772(1996)

Non Patent Literature 9: Hum. Mol. Genet. 6: 33-39, 1997

## SUMMARY OF INVENTION

### Technical Problem

An object of the present invention is to produce an antibody which targets CAPRIN-1 specifically expressed on the surface of cancer cells and has better antitumor activity than conventional antibodies, and provide use of the antibody as a therapeutic and/or preventive agent for cancer.

### Solution to Problem

The present invention has the following aspects:

The present invention provides an antibody or a fragment thereof which has immunological reactivity with a CAPRIN-1 protein, the antibody or the fragment thereof comprising a heavy chain variable region comprising amino acid sequences represented by SEQ ID NOs: 5, 6, and 7 and a light chain variable region comprising amino acid sequences represented by SEQ ID NOs: 9, 10, and 11.

In another embodiment, the antibody is a human antibody, a humanized antibody, a chimeric antibody, a single-chain antibody, or a bispecific antibody.

The present invention provides a pharmaceutical composition for treatment and/or prevention of cancer, comprising the antibody or a fragment thereof as an active ingredient.

In an embodiment of the present invention, the cancer is breast cancer, kidney cancer, pancreatic cancer, large bowel cancer, lung cancer, brain tumor, gastric cancer, uterine cervix cancer, ovary cancer, prostate cancer, urinary bladder cancer, esophageal cancer, leukemia, lymphoma, fibrosarcoma, mastocytoma, or melanoma.

The present specification encompasses the contents disclosed in Japanese Patent Application No. 2011-171377 based on which the priority of the present application is claimed.

#### Advantageous Effects of Invention

The antibody against CAPRIN-1 used in the present invention damages cancer cells. Thus, the antibody against CAPRIN-1 is useful in the treatment and/or prevention of cancer.

#### DESCRIPTION OF EMBODIMENTS

The antibody against a CAPRIN-1 polypeptide used in the present invention can be examined for its antitumor activity, as described later, by examining *in vivo* the inhibition of tumor proliferation in a cancer-bearing animal or by examining *ex vivo* the presence or absence of immunocyte- or complement-mediated cytotoxic activity exhibited by the antibody against tumor cells expressing the polypeptide.

The antibody against CAPRIN-1 used in the present invention is a monoclonal antibody specifically binding to the CAPRIN-1 protein. Such a monoclonal antibody may be any type of antibody that can exert antitumor activity and includes, for example, non-human animal antibodies (e.g., mouse, rat, rabbit, and chicken monoclonal antibodies), recombinant antibodies (e.g., synthetic antibodies, multispecific antibodies, humanized antibodies, chimeric antibodies, and single-chain antibodies (scFv)), human antibodies, and their antibody fragments (e.g., Fab, F(ab')<sub>2</sub>, and Fv). These antibodies and fragments thereof can be prepared by methods generally known to those skilled in the art. In the case of a human test subject, a human antibody or a humanized antibody is desirable for avoiding or suppressing rejection.

The antibody is any class of immunoglobulin molecule, for example, IgG, IgE, IgM, IgA, IgD, or IgY, or any subclass, for example, IgG1, IgG2, IgG3, IgG4, IgA1, or IgA2.

The antibody may be further modified by acetylation, formylation, amidation, phosphorylation, PEGylation, or the like, in addition to glycosylation.

In this context, the phrase "specifically binding to the CAPRIN-1 protein" means that the antibody specifically binds to the CAPRIN-1 protein without substantially binding to other proteins.

The test subject to receive the treatment and/or prevention of cancer according to the present invention is a mammal such as a human, a pet animal, livestock, or a sport animal, preferably a human.

Hereinafter, antigen preparation, antibody preparation, and a pharmaceutical composition according to the present invention will be described.

#### <Preparation of Antigen for Antibody Preparation>

Proteins or fragments thereof used as sensitizing antigens for obtaining the antibody against CAPRIN-1 used in the present invention are not limited by animal types serving as their origins, including humans, dogs, cattle, horses, mice, rats, and chickens. The proteins or the fragments thereof, however, are preferably selected in view of compatibility with parent cells for use in cell fusion. In general, mammal-derived proteins are preferred. Particularly, human-derived proteins are preferred. For example, when CAPRIN-1 is human CAPRIN-1, human CAPRIN-1 proteins, partial peptides thereof, or cells expressing human CAPRIN-1 can be used as sensitizing antigens.

The nucleotide sequences and amino acid sequences of human CAPRIN-1 and homologs thereof can be obtained, for

example, by making an access to GenBank (NCBI, USA) and using BLAST or FASTA algorithm (Karlin and Altschul, Proc. Natl. Acad. Sci. USA, 90: 5873-5877, 1993; and Altschul et al., Nucleic Acids Res. 25: 3389-3402, 1997).

In the present invention, with reference to the nucleotide sequence (SEQ ID NO: 1 or 3) or amino acid sequence (SEQ ID NO: 2 or 4) of human CAPRIN-1, the targets are nucleic acids or proteins consisting of sequences having 70% to 100%, preferably 80% to 100%, more preferably 90% to 100%, further preferably 95% to 100%, for example, 97% to 100%, 98% to 100%, 99% to 100%, or 99.5% to 100% sequence identity to the nucleotide sequence or amino acid sequence of the ORF or mature portion of the reference. In this context, the term "% sequence identity" means a percentage (%) of the number of identical amino acids (or bases) to the total number of amino acids (or bases) when two sequences are aligned such that the maximum degree of similarity or identity can be achieved with or without introduced gaps.

The fragments of each CAPRIN-1 protein have lengths ranging from the amino acid length of an epitope (or an antigenic determinant), which is the smallest unit recognized by the antibody, to less than the full-length of the protein. The epitope refers to a polypeptide fragment having antigenicity or immunogenicity in mammals, preferably humans. Its smallest unit consists of approximately 7 to 12 amino acid, for example, 8 to 11 amino acid.

Polypeptides comprising the above human CAPRIN-1 proteins and partial peptides thereof can be synthesized according to chemical synthesis methods, for example, Fmoc (fluorenylmethyloxycarbonyl) and tBoc (t-butyloxycarbonyl) methods (Seikagaku Jikken Koza (Biochemical Experimentation Course in English) 1, the Japanese Biochemical Society ed., Protein Chemistry IV, Chemical Modification and Peptide Synthesis, Tokyo Kagaku Dojin Co., Ltd. (Japan), 1981). Also, these polypeptides can be synthesized by routine methods using various commercially available peptide synthesizers. Alternatively, polynucleotides encoding the polypeptides may be prepared using genetic engineering approaches known in the art (Sambrook et al., Molecular Cloning, the 2nd edition, Current Protocols in Molecular Biology (1989), Cold Spring Harbor Laboratory Press; Ausubel et al., Short Protocols in Molecular Biology, the 3rd edition, A compendium of Methods from Current Protocols in Molecular Biology (1995), John Wiley & Sons; etc.) and incorporated into expression vectors, which are then introduced into host cells so that the host cells produce the polypeptides. In this way, the polypeptides of interest can be obtained.

The polynucleotides encoding the polypeptides can be readily prepared by genetic engineering approaches known in the art or routine methods using commercially available nucleic acid synthesizers. For example, a DNA comprising the nucleotide sequence of human CAPRIN-1 gene can be prepared by PCR using a human chromosomal DNA or cDNA library as a template and a pair of primers designed so as to be capable of amplifying the nucleotide sequence described in SEQ ID NO: 1. Reaction conditions for this PCR can be appropriately determined. Examples of the conditions can include, but not limited to, 30 cycles each involving reaction steps consisting of 94° C. for 30 seconds (denaturation), 55° C. for 30 seconds to 1 minute (annealing), and 72° C. for 2 minutes (elongation) using thermotolerance DNA polymerase (e.g., Taq polymerase, Pfu polymerase) and a Mg<sup>2+</sup>-containing PCR buffer, followed by reaction at 72° C. for 7 minutes. The PCR approach, conditions, etc. are described in, for example, Ausubel et al., Short Protocols in

Molecular Biology, the 3rd edition, A Compendium of Methods from Current Protocols in Molecular Biology (1995), John Wiley & Sons (particularly, Chapter 15).

Also, appropriate probes or primers can be prepared on the basis of information about the nucleotide sequences of CAPRIN-1 gene and the amino acid sequences of CAPRIN-1 proteins, and used in the screening of, for example, a human cDNA library, to isolate the desired DNA. Preferably, such a cDNA library is produced from cells, organs, or tissues expressing CAPRIN-1 proteins. Examples of such cells or tissues include cells or tissues derived from the testis or from cancers or tumors such as leukemia, breast cancer, lymphoma, brain tumor, lung cancer, pancreatic cancer, and large bowel cancer. These operations, including the preparation of probes or primers, the construction of a cDNA library, the screening of the cDNA library, and the cloning of the gene of interest, are known to those skilled in the art and can be performed according to methods described in, for example, Sambrook et al., Molecular Cloning, the 2nd edition, Current Protocols in Molecular Biology (1989), and Ausubel et al. (ibid.). DNAs encoding the human CAPRIN-1 proteins and the partial peptides thereof can be obtained from the DNA thus obtained.

The host cells may be any cell capable of expressing the above polypeptides. Examples of prokaryotic cells include, but not limited to, *E. coli*. Examples of eukaryotic cells include, but not limited to: mammalian cells such as monkey kidney cells COS 1 and Chinese hamster ovary cells CHO; a human embryonic kidney cell line HEK293; mouse embryonic skin cell line NIH3T3; yeast cells such as budding yeast and fission yeast cells; silkworm cells; and *Xenopus* egg cells.

In the case of using prokaryotic cells as the host cells, the expression vectors used have an origin that permits replication in the prokaryotic cells, a promoter, a ribosomal binding site, a multicloning site, a terminator, a drug resistance gene, an auxotrophic complementary gene, etc. Examples of expression vectors for *E. coli* can include pUC series, pBlue-script II, pET expression systems, and pGEX expression systems. The DNAs encoding the above polypeptides can be incorporated into such expression vectors, with which prokaryotic host cells are then transformed, followed by culture of the obtained transformants so that the polypeptides encoded by the DNAs are expressed in the prokaryotic host cells. In this respect, the polypeptides may be expressed as fusion proteins with other proteins.

In the case of using eukaryotic cells as the host cells, expression vectors for eukaryotic cells having a promoter, a splicing region, a poly(A) addition site, etc. are used as the expression vectors. Examples of such expression vectors can include pKA1, pCDM8, pSVK3, pMSG, pSVL, pBK-CMV, pBK-RSV, EBV, pRS, pcDNA3, and pYES2 vectors. In the same way as above, the DNAs encoding the above polypeptides can be incorporated into such expression vectors, with which eukaryotic host cells are then transformed, followed by culture of the obtained transformants so that the polypeptides encoded by the DNAs are expressed in the eukaryotic host cells. In the case of using expression vectors such as pIND/V5-His, pFLAG-CMV-2, pEGFP-N1, or pEGFP-C1, the polypeptides may be expressed as various fusion proteins tagged with His tag (e.g., (His)<sub>6</sub>-(His)<sub>10</sub>), FLAG tag, myc tag, HA tag, GFP, or the like.

The expression vectors can be introduced into the host cells using well known methods such as electroporation, a calcium phosphate method, a liposome method, a DEAE dextran method, microinjection, viral infection, lipofection, and binding with cell-penetrating peptides.

The polypeptide of interest can be isolated and purified from the host cells by a combination of separation operations known in the art. Examples thereof include, but not limited to, treatment with a denaturant (e.g., urea) or a detergent, ultrasonication, enzymatic digestion, salting-out, solvent fractionation and precipitation, dialysis, centrifugation, ultrafiltration, gel filtration, SDS-PAGE, isoelectric focusing electrophoresis, ion-exchange chromatography, hydrophobic chromatography, affinity chromatography, and reverse-phase chromatography.

#### <Structure of Antibody>

Antibodies are usually heteromultimeric glycoproteins comprising at least two heavy chains and two light chains. The immunoglobulins other than IgM are heterotetrameric glycoproteins of approximately 150 kDa each composed of two identical light (L) chains and two identical heavy (H) chains. Typically, each light chain is connected to a heavy chain via a single covalent disulfide bond, though the number of disulfide bonds between heavy chains varies among different immunoglobulin isotypes. Each of the heavy and light chains also has an intrachain disulfide bond. Each heavy chain has a variable domain (VH region) at one end, followed by a series of constant regions. Each light chain has a variable domain (VL region) at one end and has a single constant region at the other end. The light chain constant region is aligned with the first heavy chain constant region, while the light chain variable domain is aligned with the heavy chain variable domain. Particular regions called complementarity determining regions (CDRs) in the antibody variable domains exhibit specific variability and impart binding specificity to the antibody. Portions relatively conserved in the variable regions are called framework regions (FRs). The complete heavy and light chain variable domains each comprise four FRs connected via three CDRs. These three CDRs are called CDRH1, CDRH2, and CDRH3 in this order from the N-terminus of the heavy chain. Likewise, the CDRs are called CDRL1, CDRL2, and CDRL3 in the light chain. CDRH3 is most important for the binding specificity of the antibody for its antigen. In addition, CDRs in each chain are kept close to each other by the FR regions and contribute to the formation of an antigen-binding site in the antibody, together with CDRs in the other chain. The constant regions do not directly contribute to antibody-antigen binding, but exhibit various effector functions, for example, involvement in antibody-dependent cellular cytotoxicity (ADCC), phagocytosis mediated by binding to an Fcγ receptor, half-life/clearance rate mediated by a neonatal Fc receptor (FcRn), and complement-dependent cytotoxicity (CDC) mediated by a C1q component in the complement cascade.

#### <Preparation of Antibody>

The anti-CAPRIN-1 antibody according to the present invention means an antibody having immunological reactivity with a full-length CAPRIN-1 protein or a fragment thereof.

In this context, the "immunological reactivity" means the property of the antibody binding to the CAPRIN-1 antigen *in vivo*. Via such binding, the antibody exerts the function of damaging (e.g., killing, suppressing, or regressing) tumor. Specifically, the antibody used in the present invention is not limited by its type as long as the antibody can damage tumors such as breast cancer, kidney cancer, pancreatic cancer, large bowel cancer, lung cancer, brain tumor, gastric cancer, uterine cervix cancer, ovary cancer, prostate cancer, urinary bladder cancer, esophageal cancer, leukemia, lymphoma, fibrosarcoma, mastocytoma, or melanoma as a result of binding to the CAPRIN-1 protein.

Hereinafter, preparation examples of various monoclonal antibodies will be shown.

For example, breast cancer cell lines SK-BR-3 expressing CAPRIN-1 is administered to each mouse for immunization. The spleen is extracted from this mouse. After separation of spleen cells, the cells are fused with mouse myeloma cells. Clones producing antibodies having antiproliferative effect of a cancer cell are selected from among the obtained fusion cells (hybridomas). The hybridomas producing monoclonal antibodies having antiproliferative effect of a cancer cell are isolated and cultured. The antibody of interest can be prepared by purification from the culture supernatant according to a general affinity purification method.

The monoclonal antibody-producing hybridomas may be prepared, for example, as follows: first, animals are immunized with sensitizing antigens according to a method known in the art. This immunization method generally involves intraperitoneally or subcutaneously injecting the sensitizing antigens to mammals. Specifically, the sensitizing antigens are diluted with or suspended in PBS (phosphate-buffered saline), physiological saline, or the like into an appropriate amount and then mixed, if desired, with an appropriate amount of a conventional adjuvant, for example, a complete Freund's adjuvant. After emulsification, the resulting emulsion is administered to each mammal several times every 4 to 21 days. Alternatively, an appropriate carrier may be used for the immunization with sensitizing antigens.

After confirmation of a rise in the level of the desired antibody in the serum of the mammal thus immunized, immunocytes are collected from the mammal and subjected to cell fusion. Preferred examples of the immunocytes particularly include spleen cells.

Mammalian myeloma cells are used as partner parent cells to be fused with the immunocytes. Various cell lines known in the art, for example, P3U1 (P3-X63Ag8U1), P3 (P3x63Ag8.653) (J. Immunol. (1979) 123, 1548-1550), P3x63Ag8U.1 (Current Topics in Microbiology and Immunology (1978) 81, 1-7), NS-1 (Kohler, G. and Milstein, C. Eur. J. Immunol. (1976) 6, 511-519), MPC-11 (Margulies, D. H. et al., Cell (1976) 8, 405-415), SP2/0 (Shulman, M. et al., Nature (1978) 276, 269-270), FO (deSt. Groth, S. F. et al., J. Immunol. Methods (1980) 35, 1-21), S194 (Trowbridge, I. S. J. Exp. Med. (1978) 148, 313-323), and R210 (Galfre, G. et al., Nature (1979) 277, 131-133), are preferably used as the myeloma cells.

The cell fusion between the immunocytes and the myeloma cells can be performed basically according to a method known in the art, for example, the method of Kohler and Milstein (Kohler, G. and Milstein, C. Methods Enzymol. (1981) 73, 3-46).

More specifically, the cell fusion is carried out, for example, in the presence of a cell fusion promoter in a conventional nutrient medium. For example, polyethylene glycol (PEG) or hemagglutinating virus of Japan (HVJ) is used as the fusion promoter. If desired, an auxiliary such as dimethyl sulfoxide may be further added in order to enhance fusion efficiency.

The ratio between the immunocytes and the myeloma cells used can be arbitrarily set. For example, the amount of the immunocytes is preferably set to 1 to 10 times the amount of the myeloma cells. Examples of the medium that can be used in the cell fusion include RPMI 1640 and MEM media suitable for the growth of the myeloma cell lines as well as conventional media for use in this type of cell culture. In addition, a serum supplement such as fetal calf serum (FCS) may be used in combination with these media.

For the cell fusion, the immunocytes and the myeloma cells are well mixed in a predetermined amount of the medium. A PEG solution (average molecular weight: for example, approximately 1000 to 6000) preheated to approximately 37°C. is usually added to the mixture at a concentration of 30 to 60% (w/v) and mixed therewith to form the hybridomas of interest. Subsequently, procedures of sequentially adding an appropriate medium and removing the supernatant by centrifugation are repeated to remove cell fusion agents or the like unfavorable for the growth of the hybridomas.

The hybridomas thus obtained are cultured in a conventional selective medium, for example, a HAT medium (medium containing hypoxanthine, aminopterin, and thymidine) for selection. Culture in the HAT medium is continued for a period (usually, several days to several weeks) sufficient for the death of cells other than the hybridomas of interest (non-fused cells). Subsequently, hybridomas producing the antibody of interest are screened for and cloned as single clones by a conventional limiting dilution method.

In addition to such obtainment of the hybridomas by the immunization of non-human animals with antigens, hybridomas producing human antibodies having the desired activity (e.g., cell antiproliferative activity) may be obtained by sensitizing human lymphocytes, for example, EB virus-infected human lymphocytes, with proteins, protein-expressing cells, or lysates thereof in vitro and fusing the sensitized lymphocytes with human-derived myeloma cells capable of dividing permanently, for example, U266 (Accession No. TIB 196).

The monoclonal antibody-producing hybridomas thus prepared can be subcultured in a conventional medium and can also be stored for a long period in liquid nitrogen.

Specifically, the desired antigens or cells expressing the desired antigens are used as sensitizing antigens in immunization according to a conventional immunization method. The obtained immunocytes are fused with parent cells known in the art according to a conventional cell fusion method. Monoclonal antibody-producing cells (hybridomas) can be screened for by a conventional screening method to prepare the antibody of interest.

In this context, for example, KM mice (Kirin Pharma Co., Ltd./Medarex) and Xeno mice (Amgen Inc.) are known as the human antibody-producing mice (e.g., International Publication Nos. WO02/43478 and WO02/092812). Complete human polyclonal antibodies can be obtained from the blood of such mice immunized with CAPRIN-1 proteins or fragments thereof. Alternatively, spleen cells may be isolated from the mice thus immunized and fused with myeloma cells. In this way, human monoclonal antibodies can be obtained.

The antigens can be prepared according to, for example, a method using animal cells (JP Patent Publication (Kohyo) No. 2007-530068) or a method using baculovirus (e.g., International Publication No. WO98/46777). Antigens having low immunogenicity can be bound to immunogenic macromolecules such as albumin for immunization.

Alternatively, recombinant antibodies may be used, which are produced using a gene recombination technique which involves: cloning the antibody genes from hybridomas; incorporating the antibody genes into appropriate vectors; and introducing the vectors into hosts (see, e.g., Carl, A. K. Borrebaeck, James, W. Larrick, THERAPEUTIC MONOCLONAL ANTIBODIES, Published in the United Kingdom by MACMILLAN PUBLISHERS LTD, 1990). Specifically, cDNAs of the antibody variable region (V region) are synthesized from the mRNAs of hybridomas using reverse transcriptase. After obtainment of DNAs encoding the antibody V regions of interest, the DNAs are ligated with DNAs encoding the desired antibody constant regions (C regions). The result-



ing ligation products are then incorporated into expression vectors. Alternatively, the antibody V region-encoding DNAs may be incorporated into expression vectors containing antibody C region DNAs. These DNAs are incorporated into the expression vectors so as to be expressed under the control of expression control regions, for example, an enhancer and a promoter. Next, host cells can be transformed with the resulting expression vectors and allowed to express antibodies.

As described above, the anti-CAPRIN-1 antibody of the present invention is any of various monoclonal antibodies including non-human animal monoclonal antibodies, human monoclonal antibodies, and recombinant antibodies.

These non-human animal monoclonal antibodies or human monoclonal antibodies may be prepared by the culture of hybridomas obtained by the fusion between spleen cells from non-human animals (e.g., mice, human antibody-producing mice, chicken, or rabbits) immunized with CAPRIN-1 proteins and myeloma cells.

The recombinant antibody includes, as described above, chimeric antibodies, humanized antibodies, single-chain antibodies, multispecific antibodies, and the like.

The chimeric antibody can be prepared by: ligating DNAs encoding light or heavy chain variable regions of an antibody derived from a non-human animal (e.g., mouse, rat, rabbit, or chicken) with DNAs encoding light or heavy chain constant regions derived from a human antibody; incorporating the resulting fusion genes into expression vectors; and introducing the vectors into hosts so that antibodies are produced.

Specifically, the chimeric antibody is, for example, an antibody composed of variable regions of mouse antibody heavy and light chains and heavy and light chain constant region of human antibodies. The chimeric antibody can be prepared using a method known in the art which involves, for example: ligating DNAs encoding mouse antibody V regions with DNAs encoding human antibody C regions; incorporating the resulting ligation products into expression vectors; and introducing the vectors into hosts so that antibodies are produced. In Examples described later, human-mouse chimeric antibodies were prepared and confirmed to have an antitumor effect. These monoclonal antibodies comprise a heavy chain variable (VH) region having the amino acid sequence of SEQ ID NO: 8 and a light chain variable (VL) region having the amino acid sequence of SEQ ID NO: 12, wherein the VH region comprises CDR1 represented by the amino acid sequence of SEQ ID NO: 5, CDR2 represented by the amino acid sequence of SEQ ID NO: 6, and CDR3 represented by the amino acid sequence of SEQ ID NO: 7, and the VL region comprises CDR1 represented by the amino acid sequence of SEQ ID NO: 9, CDR2 represented by the amino acid sequence of SEQ ID NO: 10, and CDR3 represented by the amino acid sequence of SEQ ID NO: 11.

The humanized antibody, also called reshaped human antibody, is an engineered antibody. The humanized antibody is constructed by grafting CDRs of human antibody with CDRs corresponding to antibody derived from an immunized animal. Specifically, DNAs in which the CDR coding sequences in DNAs encoding a human antibody-derived light or heavy chain variable region are substituted by corresponding CDR coding sequences of a non-human animal (e.g., mouse, rat, rabbit, or chicken)-derived antibody are prepared and ligated with the DNAs encoding human antibody-derived light or heavy chain constant regions. The resulting fusion genes can be incorporated into expression vectors, which are then introduced into hosts for antibody production to prepare the humanized antibody of interest.

Specifically, for example, DNA sequences designed so as to link mouse and chicken antibodies CDRs and human anti-

body framework regions (FRs) are synthesized by PCR using several prepared oligonucleotides having terminal portions overlapping with each other. The obtained DNAs are ligated with DNAs encoding human antibody constant regions. Subsequently, the resulting ligation products are incorporated into expression vectors, which are then introduced into hosts for antibody production to obtain the antibody of interest (see European Patent Application Publication No. EP239400 and International Publication No. WO96/02576). The human antibody FRs connected via CDRs are selected such that the CDRs form a favorable antigen-binding site. If necessary, amino acids in the framework regions of antibody variable regions may be substituted so that the CDRs of the resulting reshaped human antibody form an appropriate antigen-binding site (Sato K. et al., Cancer Research 1993, 53: 851-856). In addition, these framework regions may be replaced with framework regions derived from various human antibodies (see International Publication No. WO99/51743).

The human antibody framework regions connected via CDRs are selected such that the CDRs form a favorable antigen-binding site. If necessary, amino acids in the framework regions of antibody variable regions may be substituted such that the CDRs of the resulting reshaped human antibody form an appropriate antigen-binding site (Sato K. et al., Cancer Research 1993, 53: 851-856).

The single-chain antibody refers to an antibody or the fragment thereof comprising variable regions of heavy and light chains linearly linked to each other via a linker. A DNA encoding the single-chain antibody can be prepared by ligating a DNA encoding the heavy chain variable region, a DNA encoding the linker, and a DNA encoding the light chain variable region. In this context, the variable regions of heavy and light chains are both derived from a human antibody or derived from a human antibody having CDRs alone substituted by CDRs of a non-human animal (e.g., mouse, rat, rabbit, or chicken)-derived antibody. The linker consists of 12 to 19 amino acids. Examples thereof include  $(G_4S)_3$  consisting of 15 amino acids (G. B. Kim et al., Protein Engineering Design and Selection 2007, 20 (9): 425-432).

The multispecific antibody refers to an antibody capable of specifically binding a plurality of different epitopes. For example, the bispecific antibody (diabody) is capable of specifically binding to two different epitopes. A DNA encoding the bispecific antibody can be prepared by ligating, for example, a DNA encoding a heavy chain variable region A, a DNA encoding a light chain variable region B, a DNA encoding a heavy chain variable region B, and a DNA encoding a light chain variable region A in this order (provided that the DNA encoding a light chain variable region B and the DNA encoding a heavy chain variable region B are ligated via a DNA encoding a linker as described above). In this context, the variable regions of heavy and light chains are all derived from a human antibody or derived from a human antibody having CDRs alone substituted by CDRs of a non-human animal (e.g., mouse, rat, rabbit, or chicken)-derived antibody.

Amino acids in variable regions (e.g., FRs) or constant regions of the chimeric antibody or the humanized antibody thus prepared may be substituted by other amino acids.

The amino acid substitution is the substitution of, for example, less than 15, less than 10, 8 or less, 7 or less, 6 or less, 5 or less, 4 or less, 3 or less, or 2 or less amino acids, preferably 1 to 5 amino acids, more preferably 1 or 2 amino acids. The substituted antibody should be functionally equivalent to an unsubstituted antibody. The substitution is desirably conservative amino acid substitution, which is the substitution between amino acids similar in properties such as charge, side chains, polarity, and aromaticity. The amino

acids can be classified in terms of similar properties into, for example: basic amino acids (arginine, lysine, and histidine); acidic amino acids (aspartic acid and glutamic acid); uncharged polar amino acids (glycine, asparagine, glutamine, serine, threonine, cysteine, and tyrosine); nonpolar amino acids (leucine, isoleucine, alanine, valine, proline, phenylalanine, tryptophan, and methionine); branched amino acids (leucine, valine, and isoleucine); and aromatic amino acids (phenylalanine, tyrosine, tryptophan, and histidine).

Examples of modified antibodies can include antibodies bound with various molecules such as polyethylene glycol (PEG). In the modified antibody of the present invention, the substance to be bound is not limited. In order to obtain such a modified antibody, the obtained antibody can be chemically modified. A method therefor has already been established in the art.

In this context, the phrase "functionally equivalent" means that an antibody concerned has biological or biochemical activity similar to that of the antibody of the present invention, specifically, the antibody concerned has the function of damaging tumor and essentially causes no rejection when applied to humans, for example. Examples of such activity can include antiproliferative activity and binding activity.

A method for preparing a polypeptide functionally equivalent to a certain polypeptide, which involves introducing a mutation into a polypeptide, is well known to those skilled in the art. For example, those skilled in the art can appropriately introduce a mutation into the antibody of the present invention using site-directed mutagenesis (Hashimoto-Gotoh, T. et al., (1995) *Gene* 152: 271-275; Zoller, M. J., and Smith, M. (1983) *Methods Enzymol.*, 100: 468-500; Kramer, W. et al., (1984) *Nucleic Acids Res.*, 12: 9441-9456; Kramer, W. and Fritz, H. J., (1987) *Methods Enzymol.*, 154: 350-367; Kunkel, T. A., (1985) *Proc. Natl. Acad. Sci. USA.*, 82: 488-492; and Kunkel (1988) *Methods Enzymol.*, 85: 2763-2766) or the like, thereby prepare an antibody functionally equivalent to the antibody of the present invention.

An antibody that recognizes an epitope of a CAPRIN-1 protein recognized by each anti-CAPRIN-1 antibody described above can be obtained by a method generally known to those skilled in the art. For example, the antibody can be obtained by a method which involves determining the epitope of the CAPRIN-1 protein recognized by the anti-CAPRIN-1 antibody by a conventional method (e.g., epitope mapping) and preparing an antibody using a polypeptide having an amino acid sequence contained in the epitope as an immunogen, or a method which involves determining an epitope for an antibody prepared by a conventional method and selecting an antibody that recognizes the same epitope as that for the anti-CAPRIN-1 antibody.

The antibody of the present invention has an affinity constant  $K_a$  ( $K_{on}/K_{off}$ ) of preferably at least  $10^7$  M<sup>-1</sup>, at least  $10^8$  M<sup>-1</sup>, at least  $5 \times 10^8$  M<sup>-1</sup>, at least  $10^9$  M<sup>-1</sup>, at least  $5 \times 10^9$  M<sup>-1</sup>, at least  $10^{10}$  M<sup>-1</sup>, at least  $5 \times 10^{10}$  M<sup>-1</sup>, at least  $10^{11}$  M<sup>-1</sup>, at least  $5 \times 10^{11}$  M<sup>-1</sup>, at least  $10^{12}$  M<sup>-1</sup>, or at least  $10^{13}$  M<sup>-1</sup>.

The antibody of the present invention can be conjugated with an antitumor agent. The conjugation of the antibody with the antitumor agent can be performed via a spacer having a group (e.g., a succinimidyl group, a formyl group, a 2-pyridyldithio group, a maleimidyl group, an alkoxycarbonyl group, or a hydroxy group) reactive with an amino group, a carboxyl group, a hydroxy group, a thiol group, or the like.

Examples of the antitumor agent include the following antitumor agents publicly known in literatures, etc.: paclitaxel, doxorubicin, daunorubicin, cyclophosphamide, methotrexate, 5-fluorouracil, thiotepa, busulfan, improsulfan, piposulfan, benzodopa, carboquone, meturedopa, uredopa,

altretamine, triethylenemelamine, triethylenephosphoramide, triethylenethiophosphoramide, trimethylolmelamine, bullatacin, bullatacinone, camptothecin, bryostatins, callistatin, cryptophycin 1, cryptophycin 8, dolastatin, duocarmycin, eleutherobin, pancratistatin, sarcodictyin, spongistatin, chlorambucil, chlornaphazine, cholophosphamide, estramustine, ifosfamide, mechlorethamine, mechlorethamine oxide hydrochloride, melphalan, novembichin, phenesterine, prednimustine, trofosfamide, uracil mustard, carmustine, chlorozotocin, fotemustine, lomustine, nimustine, ranimustine, calicheamicin, dynemicin, clodronate, esperamicin, aclacinomycin, actinomycin, authramycin, azaserine, bleomycin, cactinomycin, carabacin, carminomycin, carzinophilin, chromomycin, dactinomycin, detorbicin, 6-diazo-5-oxo-L-norleucine, Adriamycin, epirubicin, esorubicin, idarubicin, marcellomycin, mitomycin C, mycophenolic acid, nogalamycin, olivomycin, peplomycin, potfiromycin, puromycin, quelamycin, rodorubicin, streptonigrin, streptozocin, tubercidin, ubenimex, zinostatin, zorubicin, denopterin, pteropterin, trimetrexate, fludarabine, 6-mercaptopurine, thiamiprine, thioguanine, ancitabine, azacitidine, 6-azauridine, carmofur, cytarabine, dideoxyuridine, doxifluridine, enocitabine, floxuridine, androgens (e.g., calusterone, dromostanolone propionate, epitioestanol, mepitioestane, and testolactone), aminoglutethimide, mitotane, trilostane, frolinic acid, aceglutone, aldophosphamide glycoside, aminolevulinic acid, eniluracil, amsacrine, bestrabucil, bisantrene, edatraxate, defofamine, demecolcine, diaziquone, elfornithine, elliptinium acetate, epothilone, etoglucid, lentinan, lonidamine, maytansine, ansamitocin, mitoguazone, mitoxantrone, mopidanmol, nitraerine, pentostatin, phenamet, pirarubicin, losoxantrone, podophyllinic acid, 2-ethylhydrazide, procarbazine, razoxane, rhizoxin, schizophyllan, spirogermanium, tenuazonic acid, triaziquone, roridin A, anguidine, urethane, vindesine, dacarbazine, mannomustine, mitobronitol, mitolactol, pipobroman, gacytosine, docetaxel, chlorambucil, gemcitabine, 6-thioguanine, mercaptopurine, cisplatin, oxaliplatin, carboplatin, vinblastine, etoposide, ifosfamide, mitoxantrone, vincristine, vinorelbine, novantrone, teniposide, edatrexate, daunomycin, aminopterin, Xeloda, ibandronate, irinotecan, topoisomerase inhibitors, difluoromethylornithine (DMFO), retinoic acid, capecitabine, and pharmaceutically acceptable salts and derivatives thereof.

Alternatively, the antibody of the present invention can be administered in combination with an antitumor agent to produce a higher therapeutic effect. This approach is adaptable to a patient with cancer expressing CAPRIN-1 either before or after surgical operation. This approach can be applied, particularly after surgery, to CAPRIN-1-expressing cancer, which has been treated conventionally with an antitumor agent alone, to produce higher prevention of cancer recurrence or prolongation of survival time.

Examples of the antitumor agent used in the combined administration with the antibody of the present invention include the following antitumor agents publicly known in literatures, etc.: paclitaxel, doxorubicin, daunorubicin, cyclophosphamide, methotrexate, 5-fluorouracil, thiotepa, busulfan, improsulfan, piposulfan, benzodopa, carboquone, meturedopa, uredopa, altretamine, triethylenemelamine, triethylenephosphoramide, triethylenethiophosphoramide, trimethylolmelamine, bullatacin, bullatacinone, camptothecin, bryostatins, callistatin, cryptophycin 1, cryptophycin 8, dolastatin, duocarmycin, eleutherobin, pancratistatin, sarcodictyin, spongistatin, chlorambucil, chlornaphazine, cholophosphamide, estramustine, ifosfamide, mechlorethamine, mechlorethamine oxide hydrochloride, melphalan,

novembichin, phenesterine, prednimustine, trofosfamide, uracil mustard, carmustine, chlorozotocin, fotemustine, lomustine, nimustine, ranimustine, calicheamicin, dynemicin, clodronate, esperamicin, aclacinomycin, actinomycin, aithramycin, azaserine, bleomycin, cactinomycin, carabacin, carminomycin, carzinophilin, chromomycin, dactinomycin, detorbicin, 6-diazo-5-oxo-L-norleucine, Adriamycin, epirubicin, esorubicin, idarubicin, marcellomycin, mitomycin C, mycophenolic acid, nogalamycin, olivomycin, peplomycin, potfiromycin, puromycin, quelamycin, rodorubicin, streptonigrin, streptozocin, tubercidin, ubenimex, zinostatin, zorubicin, denopterin, pteropterin, trimetrexate, fludarabine, 6-mercaptapurine, thiamiprine, thioguanine, ancitabine, azacitidine, 6-azauridine, carmofer, cytarabine, dideoxyuridine, doxifluridine, enocitabine, floxuridine, calusterone, dromostanolone propionate, epitostanol, mepitiostane, testolactone, aminoglutethimide, mitotane, trilostane, frolinic acid, aceglatone, aldophosphamide glycoside, aminolevulinic acid, eniluracil, amsacrine, bestabucil, bisantrene, edatraxate, defofamine, demecolcine, diaziqune, elfornithine, elliptinium acetate, epothilone, etoglucid, lentinan, lonidamine, maytansine, ansamitocin, mitoguazone, mitoxantrone, mopidanmol, nitraerine, pentostatin, phenamet, pirarubicin, losoxantrone, podophyllinic acid, 2-ethylhydrazide, procarbazine, razoxane, rhizoxin, schizophyllan, spirogermanium, tenuazonic acid, triaziquone, roridin A, anguidine, urethane, vindesine, dacarbazine, mannomustine, mitobronitol, mitolactol, pipobroman, gacytosine, docetaxel, chlorambucil, gemcitabine, 6-thioguanine, mercaptopurine, cisplatin, oxaliplatin, carboplatin, vinblastine, etoposide, ifosfamide, mitoxantrone, vincristine, vinorelbine, novantrone, teniposide, edatrexate, daunomycin, aminopterin, Xeloda, ibandronate, irinotecan, topoisomerase inhibitors, difluoromethylornithine (DMFO), retinoic acid, capecitabine, and pharmaceutically acceptable salts (known in the art) and derivatives (known in the art) thereof. Of these antitumor agents, cyclophosphamide, paclitaxel, docetaxel, or vinorelbine is particularly preferably used.

Alternatively, the antibody of the present invention may be bound to a radioisotope publicly known in literatures, etc., such as  $^{211}\text{At}$ ,  $^{131}\text{I}$ ,  $^{125}\text{I}$ ,  $^{90}\text{Y}$ ,  $^{186}\text{Re}$ ,  $^{188}\text{Re}$ ,  $^{153}\text{Sm}$ ,  $^{212}\text{Bi}$ ,  $^{32}\text{P}$ ,  $^{175}\text{Lu}$ , or  $^{176}\text{Lu}$ . Desirably, a radioisotope effective for the treatment or diagnosis of tumor is used.

The antibody of the present invention is an antibody having immunological reactivity with CAPRIN-1, an antibody specifically recognizing CAPRIN-1, or an antibody specifically binding to CAPRIN-1 and exhibits cytotoxic activity or antiproliferative effect on cancer. The antibody preferably should have a structure that causes little or no rejection in recipient animals. Examples of such antibodies include human antibodies, humanized antibodies, chimeric antibodies (e.g., human-mouse chimeric antibodies), single-chain antibodies, and bispecific antibodies when the recipient animals are humans. These antibodies (1) have variable regions of heavy and light chains derived from a human antibody, (2) have variable regions with CDRs (CDR1, CDR2, and CDR3) of heavy and light chains derived from a non-human animal antibody and framework regions derived from a human antibody, or (3) these antibodies are recombinant antibodies having variable regions of heavy and light chains derived from a non-human animal antibody and heavy and light chain constant regions derived from a human antibody. The antibody of the present invention is preferably antibody (1) or (2).

Such recombinant antibodies can be prepared as follows: DNAs encoding monoclonal antibodies (e.g., human, mouse, rat, rabbit, and chicken monoclonal antibodies) against human CAPRIN-1 are cloned from antibody-producing cells

such as hybridomas and used as templates in RT-PCR or the like to prepare DNAs encoding the variable regions of the light and heavy chains in the antibodies. The respective sequences of the variable regions of the light and heavy chains and the respective sequences of CDR1, CDR2, and CDR3 in each region are determined on the basis of the Kabat EU numbering system (Kabat et al., Sequences of Proteins of Immunological Interest, 5th Ed. Public Health Service, National Institute of Health, Bethesda, Md. (1991)).

Such a DNA encoding each variable region or a DNA encoding each CDR is prepared using a gene recombination technique (Sambrook et al., Molecular Cloning A Laboratory Manual, Cold Spring Harbor Laboratory Press (1989)) or a DNA synthesizer. In this context, the human monoclonal antibody-producing hybridomas can be prepared by immunizing human antibody-producing animals (e.g., mice) with human CAPRIN-1 and then fusing spleen cells excised from the immunized animals with myeloma cells. Aside from this, DNAs encoding light or heavy chain variable and constant regions derived from human antibody are prepared, if necessary, using a gene recombination technique or a DNA synthesizer.

The prepared recombinant DNAs can be incorporated into one or more appropriate vectors, which are then introduced into host cells (e.g., mammalian cells, yeast cells, and insect cells) so that the DNAs are (co)expressed to produce recombinant antibodies (P. J. Delves., ANTIBODY PRODUCTION ESSENTIAL TECHNIQUES., 1997 WILEY, P. Shepherd and C. Dean., Monoclonal Antibodies., 2000 OXFORD UNIVERSITY PRESS; and J. W. Goding., Monoclonal Antibodies: principles and practice., 1993 ACADEMIC PRESS).

Examples of the antibody of the present invention prepared by any of the methods described above include the following antibody (a) obtained in Examples described later:

(a) an antibody or the fragment thereof comprising a heavy chain variable region comprising amino acid sequences represented by SEQ ID NOs: 5, 6, and 7 and a light chain variable region comprising amino acid sequences represented by SEQ ID NOs: 9, 10, and 11 (e.g., an antibody composed of a heavy chain variable region consisting of the amino acid sequence represented by SEQ ID NO: 8 and a light chain variable region consisting of the amino acid sequence represented by SEQ ID NO: 12).

In this context, the amino acid sequences represented by SEQ ID NOs: 5, 6, and 7 correspond to CDR1, CDR2, and CDR3, respectively, of a heavy chain variable region of a mouse antibody. The amino acid sequences represented by SEQ ID NOs: 9, 10, and 11 correspond to CDR1, CDR2, and CDR3, respectively, of a light chain variable region of a mouse antibody.

Examples of the humanized antibody, the chimeric antibody, the single-chain antibody, or the multispecific antibody of the present invention include the following antibodies (i), (ii), and (iii):

(i) an antibody or the fragment thereof comprising a heavy chain variable region comprising the amino acid sequences represented by SEQ ID NOs: 5, 6, and 7 and the amino acid sequences of framework regions derived from human antibody and a light chain variable region comprising the amino acid sequences represented by SEQ ID NOs: 9, 10, and 11 and the amino acid sequences of framework regions derived from human antibody;

(ii) an antibody or the fragment thereof comprising a heavy chain variable region comprising the amino acid sequences represented by SEQ ID NOs: 5, 6, and 7 and the amino acid sequences of framework regions derived from human antibody, a heavy chain constant region comprising an amino acid

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sequence derived from human antibody, a light chain variable region comprising the amino acid sequences represented by SEQ ID NOs: 9, 10, and 11 and the amino acid sequences of framework regions derived from human antibody, and a light chain constant region comprising an amino acid sequence derived from human antibody; and

(iii) an antibody or the fragment thereof comprising a heavy chain variable region comprising the amino acid sequence represented by SEQ ID NO: 8, a heavy chain constant region comprising an amino acid sequence derived from human antibody, a light chain variable region comprising the amino acid sequence represented by SEQ ID NO: 12, and a light chain constant region comprising an amino acid sequence derived from human antibody.

The sequences of the constant and variable regions of human antibody heavy and light chains are available from, for example, NCBI (USA; GenBank, UniGene, etc.). For example, the following sequences can be referred to: Accession No. J00228 for a constant region of a human IgG1 heavy chain; Accession No. J00230 for a constant region of a human IgG2 heavy chain; Accession No. X03604 for a constant region of a human IgG3 heavy chain; Accession No. K01316 for a constant region of a human IgG4 heavy chain; Accession Nos. V00557, X64135, and X64133 for a constant region of a human light chain  $\kappa$ ; and Accession Nos. X64132 and X64134 for a constant region of a human light chain  $\lambda$ .

Preferably, these antibodies have cytotoxic activity and can thereby exert an antitumor effect.

The above particular sequences of the variable regions and CDRs of heavy and light chains in each antibody are provided merely for illustrative purposes. It is obvious that the antibody of the present invention is not limited by the particular sequences. Hybridomas capable of producing anti-human CAPRIN-1 human antibodies or non-human animal antibodies (e.g., mouse antibodies) different from those described above are prepared, and monoclonal antibodies produced by the hybridomas are recovered and assessed as being (or being not) the antibodies of interest with immunological binding activity against human CAPRIN-1 and cytotoxic activity as indexes. The monoclonal antibody-producing hybridomas of interest are thereby identified. Then, DNAs encoding variable regions heavy and light chains of the in antibodies of interest are produced from the hybridomas and sequenced, as described above. The DNAs are used for the preparation of the different antibodies.

The antibody of the present invention may be the antibody (a) having the substitution, deletion, or addition of one or several amino acids, particularly in a sequence of a framework region and/or a constant region, as long as the antibody has such specificity that it can specifically recognize CAPRIN-1. In this context, the term "several" means preferably 2 to 5, more preferably 2 or 3.

The present invention further provides a DNA encoding the antibody of the present invention, a DNA encoding the heavy or light chain of the antibody, or a DNA encoding the variable region of the heavy or light chain in the antibody. Such a DNA includes, for example, a DNA encoding a heavy chain variable region which comprises nucleotide sequences encoding the amino acid sequences represented by SEQ ID NOs: 5, 6, and 7, and a DNA encoding a light chain variable region which comprises nucleotide sequences encoding the amino acid sequences represented by SEQ ID NOs: 9, 10, and 11, in the case of the antibody (a).

The CDRs encoded by the DNA having these sequences serve as regions that determine the specificity of the antibody. Sequences encoding the other regions (i.e., constant regions and framework regions) of the antibody may therefore be

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sequences derived from other antibodies. In this context, "other antibodies" also include antibodies derived from non-human organisms and are preferably those derived from humans from the viewpoint of reducing side effects. Specifically, the DNA of the present invention preferably comprises nucleotide sequences encoding corresponding human antibody-derived amino acid sequences of regions encoding each framework region and each constant region in the heavy and light chains.

Further examples of the DNA encoding the antibody of the present invention include a DNA encoding a heavy chain variable region which comprises a nucleotide sequence encoding the amino acid sequence represented by SEQ ID NO: 8, and a DNA encoding a light chain variable region which comprises a nucleotide sequence encoding the amino acid sequence represented by SEQ ID NO: 12. In this context, the nucleotide sequence encoding the amino acid sequence represented by SEQ ID NO: 8 is, for example, the nucleotide sequence represented by SEQ ID NO: 13. The nucleotide sequence encoding the amino acid sequence represented by SEQ ID NO: 12 is, for example, the nucleotide sequence represented by SEQ ID NO: 14. For these DNAs, it is preferred that a region encoding each constant region in the heavy and light chains should comprise a nucleotide sequence encoding a corresponding amino acid sequence derived from human antibody.

These antibody DNAs can be obtained, for example, by the methods described above or the following method: first, total RNAs are prepared from hybridomas related to the antibody of the present invention using a commercially available RNA extraction kit, and cDNAs are synthesized using reverse transcriptase and random primers or the like. Subsequently, the antibody-encoding cDNAs are amplified by PCR using oligonucleotide primers for conserved sequences of each variable region in known mouse antibody heavy and light chain genes. Sequences encoding the constant regions can be obtained by the PCR amplification of known sequences. The nucleotide sequence of the DNA can be incorporated into a plasmid or a phage for sequencing, for example, and determined according to a routine method.

The antitumor effect of the anti-CAPRIN-1 antibody used in the present invention on CAPRIN-1-expressing cancer cells seems to be brought about by effector cell-mediated antibody-dependent cellular cytotoxicity (ADCC) and complement-dependent cytotoxicity (CDC) against the CAPRIN-1-expressing cells.

The antitumor effect based on the mechanism is known to correlate with the number of antibody-binding target molecules expressed on the surface of cancer cells (Niwa R., Clinical Cancer Research 2005 Mar. 15; 11 (6): 2327-2336). The number of target molecules expressed on the surface of cancer cells can be examined using an existing assay kit capable of measuring the number of molecules on cell surface. Specifically, the number of antibody-binding target molecules can be determined by: reacting cancer cells with, for example, antibodies against the target molecules as primary antibodies; reacting therewith fluorescently labeled antibodies against the primary antibodies, together with calibration curve beads with the preliminarily known number of molecules; measuring the mean fluorescence intensity of the samples; and determining the number of the target molecules on the basis of the obtained calibration curve.

Thus, the anti-CAPRIN-1 antibody used in the present invention can be evaluated for its activity by determining *ex vivo* the ADCC activity or the CDC activity against CAPRIN-1-expressing cancer cells or by examining the number of CAPRIN-1 molecules expressed on the surface of cancer

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cells in the case of using the anti-CAPRIN-1 antibody according to the present invention as a primary antibody as specifically shown below in Examples.

The anti-CAPRIN-1 antibody used in the present invention binds to CAPRIN-1 proteins on cancer cells and exhibits an antitumor effect through the activity. Thus, the anti-CAPRIN-1 antibody of the present invention is presumably useful in the treatment or prevention of cancer. Specifically, the present invention provides a pharmaceutical composition for treatment and/or prevention of cancer, comprising the anti-CAPRIN-1 antibody as an active ingredient. The anti-CAPRIN-1 antibody used for the purpose of administration to human bodies (antibody therapy) is preferably a human antibody or a humanized antibody for reducing immunogenicity.

The anti-CAPRIN-1 antibody with higher binding affinity for a CAPRIN-1 protein on cancer cell surface exerts stronger antitumor activity. Thus, a stronger antitumor effect can be expected if the anti-CAPRIN-1 antibody having high binding affinity for the CAPRIN-1 protein can be obtained. Such an antibody is adaptable to a pharmaceutical composition intended for the treatment and/or prevention of cancer. Desirably, such high binding affinity is preferably at least  $10^7 \text{ M}^{-1}$ , at least  $10^8 \text{ M}^{-1}$ , at least  $5 \times 10^8 \text{ M}^{-1}$ , at least  $10^9 \text{ M}^{-1}$ , at least  $5 \times 10^9 \text{ M}^{-1}$ , at least  $10^{10} \text{ M}^{-1}$ , at least  $5 \times 10^{10} \text{ M}^{-1}$ , at least  $10^{11} \text{ M}^{-1}$ , at least  $5 \times 10^{11} \text{ M}^{-1}$ , at least  $10^{12} \text{ M}^{-1}$ , or at least  $10^{13} \text{ M}^{-1}$ , in terms of an association constant (affinity constant)  $K_a$  ( $k_{on}/k_{off}$ ), as described above.

The binding of anti-CAPRIN-1 antibodies to a larger number of CAPRIN-1 molecules on cancer cell surface produces stronger antitumor activity. Desirably, the number of CAPRIN-1 molecules to which the antibodies bind for the expected antitumor effect is  $10^4$  or more, preferably  $10^5$  or more CAPRIN-1 molecules per cancer cell measured using the anti-CAPRIN-1 antibody of the present invention.

#### <Binding to Antigen-Expressing Cells>

The ability of the antibody to bind to CAPRIN-1 can be determined by use of binding assay using, for example, ELISA, Western blot, immunofluorescence, and flow cytometry analysis, as described in Examples.

#### <Immunohistochemical Staining>

The antibody that recognizes CAPRIN-1 can be tested for its reactivity with CAPRIN-1 by an immunohistochemical method well known to those skilled in the art using a paraformaldehyde- or acetone-fixed frozen section or paraformaldehyde-fixed paraffin-embedded section of a tissue obtained from a patient during surgical operation or from an animal carrying a xenograft tissue inoculated with a cell line expressing CAPRIN-1 either spontaneously or after transfection.

For immunohistochemical staining, the antibody reactive with CAPRIN-1 can be stained by various methods. For example, the antibody can be visualized through reaction with a horseradish peroxidase-conjugated goat anti-mouse antibody or goat anti-chicken antibody.

#### <Pharmaceutical Composition>

A target of the pharmaceutical composition for treatment and/or prevention of cancer of the present invention is not particularly limited as long as the target is cancer (cells) expressing a CAPRIN-1 gene.

The terms "tumor" and "cancer" used herein mean malignant neoplasm and are used interchangeably with each other.

The cancer targeted in the present invention is cancer expressing a CAPRIN-1 protein-encoding gene and is preferably breast cancer, kidney cancer, pancreatic cancer, large bowel cancer, lung cancer, brain tumor, gastric cancer, uterine cervix cancer, ovary cancer, prostate cancer, urinary bladder

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cancer, esophageal cancer, leukemia, lymphoma, fibrosarcoma, mastocytoma, or melanoma.

Specific examples of these cancers include, but not limited to, breast adenocarcinoma, complex-type breast adenocarcinoma, malignant mixed tumor of mammary gland, intraductal papillary adenocarcinoma, lung adenocarcinoma, squamous cell cancer, small-cell cancer, large-cell cancer, glioma which is tumor of neuroepithelial tissue, ependymoma, neuronal tumor, embryonal neuroectodermal tumor, neurilemmoma, neurofibroma, meningioma, chronic lymphocytic leukemia, lymphoma, gastrointestinal lymphoma, alimentary lymphoma, small to medium cell-type lymphoma, cecal cancer, ascending colon cancer, descending colon cancer, transverse colon cancer, sigmoid colon cancer, rectal cancer, epithelial ovarian cancer, germ cell tumor, stromal cell tumor, pancreatic ductal carcinoma, invasive pancreatic ductal carcinoma, pancreatic adenocarcinoma, acinar cell carcinoma, adenosquamous carcinoma, giant cell tumor, intraductal papillary-mucinous neoplasm, mucinous cystic neoplasm, pancreatoblastoma, serous cystadenocarcinoma, solid-pseudopapillary tumor, gastrinoma, glucagonoma, insulinoma, multiple endocrine neoplasia type-1 (Wermer's syndrome), nonfunctional islet cell tumor, somatostatinoma, and VIPoma.

The test subject as the recipient are preferably mammals, for example, mammals including primates, pet animals, livestock, and sport animals and are particularly preferably humans, dogs, and cats.

In the case of using the antibody of the present invention as a pharmaceutical composition, the pharmaceutical composition can be formulated by a method generally known to those skilled in the art. For example, the pharmaceutical composition can be used in the form of a parenteral injection of an aseptic solution or suspension with water or any other pharmaceutically acceptable liquid. For example, the pharmaceutical composition may be formulated with the antibody mixed in a unit dosage form required for generally accepted pharmaceutical practice, in appropriate combination with pharmacologically acceptable carriers or media, specifically, sterilized water, physiological saline, plant oil, an emulsifier, a suspending agent, a detergent, a stabilizer, a flavoring agent, an excipient, a vehicle, a preservative, a binder, etc. The amount of the active ingredient in such a preparation is determined so that an appropriate dose within the prescribed range can be achieved.

An aseptic composition for injection can be formulated according to conventional pharmaceutical practice using a vehicle such as injectable distilled water.

Examples of aqueous solutions for injection include physiological saline, isotonic solutions containing glucose and other adjuvants, for example, D-sorbitol, D-mannose, D-mannitol, and sodium chloride. These solutions may be used in combination with an appropriate solubilizer, for example, an alcohol (specifically, ethanol) or a polyalcohol (e.g., propylene glycol and polyethylene glycol), or a non-ionic detergent, for example, polysorbate 80™ or HCO-60.

Examples of oily solutions include sesame oil and soybean oil. These solutions may be used in combination with a solubilizer such as benzyl benzoate or benzyl alcohol. The solutions may be further mixed with a buffer (e.g., a phosphate buffer solution and a sodium acetate buffer solution), a soothing agent (e.g., procaine hydrochloride), a stabilizer (e.g., benzyl alcohol and phenol), and an antioxidant. The injection solutions thus prepared are usually charged into appropriate ampules.

The pharmaceutical composition of the present invention is administered orally or parenterally, preferably parenterally.

Specific examples of its dosage forms include injections, intranasal administration agents, transpulmonary administration agents, and percutaneous administration agents. Examples of the injections include intravenous injection, intramuscular injection, intraperitoneal injection, and subcutaneous injection, through which the pharmaceutical composition can be administered systemically or locally.

Also, the administration method can be appropriately selected depending on the age, weight, sex, symptoms, etc. of a patient. The dose of a pharmaceutical composition containing the antibody or a polynucleotide encoding the antibody can be selected within a range of, for example, 0.0001 to 1000 mg/kg of body weight per dose. Alternatively, the dose can be selected within a range of, for example, 0.001 to 100000 mg/body of a patient, though the dose is not necessarily limited to these numeric values. Although the dose and the administration method vary depending on the weight, age, sex, symptoms, etc. of a patient, those skilled in the art can appropriately select the dose and the method.

The pharmaceutical composition including the antibody or fragments thereof of the present invention can be administered to a test subject to treat and/or prevent cancer, preferably breast cancer, kidney cancer, pancreatic cancer, large bowel cancer, lung cancer, brain tumor, gastric cancer, uterine cervix cancer, ovary cancer, prostate cancer, urinary bladder cancer, esophageal cancer, leukemia, lymphoma, fibrosarcoma, mastocytoma, or melanoma.

The present invention further encompasses a method for treating and/or preventing cancer, comprising administering the pharmaceutical composition of the present invention in combination with the antitumor agent as exemplified above or a pharmaceutical composition comprising the antitumor agent to a test subject. The antibody or the fragment thereof of the present invention may be administered simultaneously with or separately from the antitumor agent to the test subject. In the case of separately administering these pharmaceutical compositions, either one may be administered first or later. Their dosing intervals, doses, administration routes, and the number of doses can be appropriately selected by a specialist. The dosage forms of separate drugs to be administered simultaneously also include, for example, pharmaceutical compositions each formulated by mixing the antibody or the fragment thereof of the present invention and the antitumor agent into a pharmacologically acceptable carrier (or medium). The above descriptions about prescription, formulation, administration routes, doses, cancer, etc. as to the pharmaceutical compositions and dosage forms containing the antibody of the present invention are also applicable to any of the above-described pharmaceutical compositions and dosage forms containing the antitumor agent.

Thus, the present invention also provides a pharmaceutical combination for treatment and/or prevention of cancer, comprising the pharmaceutical composition of the present invention and a pharmaceutical composition comprising the antitumor agent as exemplified above. The present invention also provides a pharmaceutical composition for treatment and/or prevention of cancer, comprising the antibody or the fragment thereof of the present invention and the antitumor agent together with a pharmacologically acceptable carrier.  
<Polypeptide and DNA>

The present invention further provides the following polypeptides and DNAs related to the antibody (a):

(i) a polypeptide comprising the amino acid sequences represented by SEQ ID NOs: 8 and 12, and a DNA encoding the polypeptide, for example, a DNA comprising the nucleotide sequences represented by SEQ ID NOs: 13 and 14;

(ii) a heavy chain CDR polypeptide selected from the group consisting of amino acid sequences represented by SEQ ID NOs: 5, 6, and 7, and a DNA encoding the polypeptide; and

(iii) a light chain CDR polypeptide selected from the group consisting of amino acid sequences represented by SEQ ID NOs: 9, 10, and 11, and a DNA encoding the polypeptide.

These polypeptides and DNAs can be prepared using gene recombination techniques as described above.

#### <SUMMARY OF THE PRESENT INVENTION>

The aspects of the present invention described above are summarized below.

(1) An antibody or a fragment thereof which has immunological reactivity with a CAPRIN-1 protein, the antibody or the fragment thereof comprising a heavy chain variable region comprising CDRs of SEQ ID NOs: 5, 6, and 7 and a light chain variable region comprising CDRs of SEQ ID NOs: 9, 10, and 11.

(2) The antibody or the fragment thereof according to (1), wherein the antibody is a human antibody, a humanized antibody, a chimeric antibody, a single-chain antibody, or a bispecific antibody.

(3) The antibody or the fragment thereof according to (1) or (2), wherein the antibody or the fragment thereof is conjugated with an antitumor agent.

(4) A pharmaceutical composition for treatment and/or prevention of cancer comprising an antibody or the fragment thereof as an active ingredient, wherein the antibody comprises a heavy chain variable region comprising CDRs of SEQ ID NOs: 5, 6, and 7 and a light chain variable region comprising CDRs of SEQ ID NOs: 9, 10, and 11 and has immunological reactivity with the CAPRIN-1 protein.

(5) The pharmaceutical composition according to (4), wherein the cancer is breast cancer, kidney cancer, pancreatic cancer, large bowel cancer, lung cancer, brain tumor, gastric cancer, uterine cervix cancer, ovary cancer, prostate cancer, urinary bladder cancer, esophageal cancer, leukemia, lymphoma, fibrosarcoma, mastocytoma, or melanoma.

(6) A pharmaceutical combination for treatment and/or prevention of cancer, comprising a pharmaceutical composition according to (4) or (5) and a pharmaceutical composition comprising an antitumor agent.

(7) A DNA encoding an antibody or a fragment thereof according to (1) or (2).

(8) A method for treating and/or preventing cancer, comprising administering an antibody or a fragment thereof according to any of (1) to (3), a pharmaceutical composition according to (4) or (5), or a pharmaceutical combination according to (6) to a test subject.

#### EXAMPLES

Hereinafter, the present invention will be described specifically with reference to Examples. However, the scope of the present invention is not intended to be limited by these specific examples.

##### Example 1

##### Analysis of CAPRIN-1 Gene Expression in Each Tissue

Gene expression of CAPRIN-1 in canine and human normal tissues and various cell lines was examined by RT-PCR according to Example 1(4) described in WO2010/016526. As a result, strong expression was observed in the testis among the healthy canine tissues, whereas expression was observed

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in canine breast cancer and adenocarcinoma tissues. As a result of also confirming the expression in human tissues, the expression was confirmed only in the testis among normal tissues, as with the canine CAPRIN-1 gene. By contrast, the expression was detected in many types of cancer cell lines, including 8 human breast cancer cell lines (ZR75-1, MCF7, T47D, SK-BR-3, MDA-MB-157, BT-20, MDA-MB-231V, and MRK-nu-1) and 4 pancreatic cancer cell lines (Capan-2, MIAPaCa-2, Panc-1, and BxPc-3), among cancer cells. These results demonstrated that CAPRIN-1 is expressed in various cancer cells, though its expression is not seen in normal tissues other than the testis.

## Example 2

## Preparation of Mouse Monoclonal Antibody Against CAPRIN-1

100 µg of human CAPRIN-1 proteins having the amino acid sequence of SEQ ID NO: 2, which were prepared in Example 3 described in WO2010/016526, was mixed with an equal amount of MPL+TDM adjuvant (Sigma-Aldrich Corp.). This mixture was used as an antigen solution per mouse. The antigen solution was intraperitoneally administered to each 6-week-old Balb/c mouse (Japan SLC, Inc.). Then, 7 boosters were performed every 1 week to complete immunization. Three days after the final shot, the spleen of each mouse was excised and ground between two sterilized glass slides. Procedures of washing with PBS(-) (Nissui Pharmaceutical Co., Ltd.) and removing the supernatant by centrifugation at 1500 rpm for 10 minutes were repeated three times to obtain spleen cells. The obtained spleen cells were mixed with mouse myeloma cells SP2/0 (purchased from ATCC) at a ratio of 10:1. 200 µl of an RPMI1640 medium containing 10% FBS was heated to 37° C. and mixed with 800 of PEG1500 (Boehringer Ingelheim GmbH), and the PEG solution thus prepared was added to the cell mixture, which was then left standing for 5 minutes for cell fusion. After removal of the supernatant by centrifugation at 1700 rpm for 5 minutes, the cells were suspended in 150 ml of an RPMI1640 medium containing 15% FBS supplemented with 2% equivalent of a HAT solution (Life Technologies, Inc./Gibco) (HAT selective medium). This suspension was inoculated to fifteen 96-well plates (Thermo Fisher Scientific Inc./Nunc) at a concentration of 100 µl/well. The spleen cells and the myeloma cells were fused by culture at 37° C. for 7 days under conditions of 5% CO<sub>2</sub> to obtain hybridomas.

The prepared hybridomas were screened with the binding affinity of antibodies produced by the hybridomas against CAPRIN-1 proteins as an index. One µg/ml solution of the CAPRIN-1 proteins prepared in Example 3 described in WO2010/016526 was added to a 96-well plate at a concentration of 100 µl/well and left standing at 4° C. for 18 hours. Each well was washed three times with PBS-T. Then, a 0.5% bovine serum albumin (BSA) solution (Sigma-Aldrich Corp.) was added thereto at a concentration of 400 µl/well and left standing at room temperature for 3 hours. The solution in each well was discarded, and each well was washed three times with 400 µl of PBS-T. Then, the culture supernatant of each hybridoma obtained above was added thereto at a concentration of 100 µl/well and left standing at room temperature for 2 hours. Each well was washed three times with PBS-T. Then, HRP-labeled anti-mouse IgG (H+L) antibodies (Invitrogen Corp.) diluted 5000-fold with PBS were added thereto at a concentration of 100 µl/well and left standing at room temperature for 1 hour. Each well was washed three times with PBS-T. Then, a TMB substrate solution (Thermo Fisher Sci-

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entific Inc.) was added thereto at a concentration of 100 µl/well and left standing for 15 to 30 minutes to cause color reaction. After the color development, the reaction was terminated by the addition of 1 N sulfuric acid at a concentration of 100 µl/well. The absorbance was measured at 450 nm and 595 nm using an absorption spectrometer. As a result, several hybridomas producing antibodies having high absorbance were selected.

The selected hybridomas were added to a 96-well plate at a density of 0.5 cells/well and cultured in the plate. One week later, hybridomas forming single colonies in the wells were observed. The cells in these wells were further cultured, and the cloned hybridomas were screened with the binding affinity of antibodies produced by the hybridomas against CAPRIN-1 proteins as an index. One µg/ml solution of the CAPRIN-1 proteins prepared in Example 3 described in WO2010/016526 was added to a 96-well plate at a concentration of 100 µl/well and left standing at 4° C. for 18 hours. Each well was washed three times with PBS-T. Then, a 0.5% BSA solution was added thereto at a concentration of 400 µl/well and left standing at room temperature for 3 hours. The solution in each well was discarded, and each well was washed three times with 400 µl of PBS-T. Then, the culture supernatant of each hybridoma obtained above was added thereto at a concentration of 100 µl/well and left standing at room temperature for 2 hours. Each well was washed three times with PBS-T. Then, HRP-labeled anti-mouse IgG (H+L) antibodies (Invitrogen Corp.) diluted 5000-fold with PBS were added thereto at a concentration of 100 µl/well and left standing at room temperature for 1 hour. Each well was washed three times with PBS-T. Then, a TMB substrate solution (Thermo Fisher Scientific Inc.) was added thereto at a concentration of 100 µl/well and left standing for 15 to 30 minutes to cause color reaction. After the color development, the reaction was terminated by the addition of 1 N sulfuric acid at a concentration of 100 µl/well. The absorbance was measured at 450 nm and 595 nm using an absorption spectrometer. As a result, 112 hybridoma lines producing monoclonal antibodies reactive with CAPRIN-1 proteins were obtained.

Next, these monoclonal antibodies were screened for antibodies reactive with the surface of breast cancer cells expressing CAPRIN-1. Specifically, 10<sup>6</sup> cells of a human breast cancer cell line MDA-MB-231V were centrifuged in a 1.5-mL microcentrifuge tube. 100 µL of the culture supernatant of each hybridoma obtained above was added thereto and left standing for 1 hour on ice. After washing with PBS, FITC-labeled goat anti-mouse IgG antibodies (Invitrogen Corp.) diluted 500-fold with PBS containing 0.1% FBS were added thereto and left standing for 1 hour on ice. After washing with PBS, the fluorescence intensity was measured using FACS Calibur (Becton, Dickinson and Company). On the other hand, the same operation as above was performed using the serum of each untreated 6-week-old Balb/c mouse diluted 500-fold with a medium for hybridoma culture, instead of the antibodies, to prepare a control. As a result, one monoclonal antibody (#1) having stronger fluorescence intensity than that of the control, i.e., reactive with the surface of breast cancer cells, was selected.

## Example 3

## Characterization of Selected Monoclonal Antibody

The monoclonal antibodies obtained in Example 2 were analyzed according to a method described in Example 5 of WO2010/016526 for their nucleotide sequences and amino



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acid sequences encoded thereby. The resulting monoclonal antibody #1 was composed of a heavy chain variable region consisting of the amino acid sequence represented by SEQ ID NO: 8 and a light chain variable region consisting of the amino acid sequence represented by SEQ ID NO: 12. The resulting nucleotide sequence encoding the heavy chain variable region of the monoclonal antibody #1 is shown in SEQ ID NO: 13, and the amino acid sequence is shown in SEQ ID NO: 8. The nucleotide sequence encoding the light chain variable region thereof is shown in SEQ ID NO: 14, and the amino acid sequence is shown in SEQ ID NO: 12.

Specifically, the monoclonal antibody #1 was confirmed to consist of the heavy chain variable region consisting of the amino acid sequence represented by SEQ ID NO: 8 and the light chain variable region consisting of the amino acid sequence represented by SEQ ID NO: 12, wherein the heavy chain variable region had CDR1, CDR2, and CDR3 consisting of amino acid sequences of SEQ ID NOs: 5, 6, and 7, respectively, and the light chain variable region had CDR1, CDR2, and CDR3 consisting of amino acid sequences of SEQ ID NOs: 9, 10, and 11, respectively.

## Example 4

## Preparation of Human-Mouse Chimeric Monoclonal Antibody

Both ends of the gene amplification fragment comprising the nucleotide encoding the heavy chain variable region of the mouse monoclonal antibody #1 obtained in Example 3, which consists of a nucleotide sequence represented by SEQ ID NO: 13, was treated with restriction enzymes, then purified, and inserted according to a routine method into a pcDNA4/myc-His vector (Invitrogen Corp.) already having gene inserts of a mouse antibody-derived leader sequence and a human IgG<sub>1</sub> H chain constant region comprising SEQ ID NO: 37. Also, the gene amplification fragment comprising the gene of the light chain variable region of the mouse monoclonal antibody #1 consisting of a nucleotide sequence represented by SEQ ID NO: 14 was treated at both ends with restriction enzymes, then purified, and inserted according to a routine method into a pcDNA3.1/myc-His (Invitrogen Corp.) vector already having gene inserts of a mouse antibody-derived leader sequence and a human IgG<sub>1</sub> L chain constant region comprising SEQ ID NO: 38.

Next, the recombinant vector having the insert of the heavy chain variable region (SEQ ID NO: 13) of the mouse monoclonal antibody #1 and the recombinant vector having the insert of the light chain variable region (SEQ ID NO: 14) of the mouse monoclonal antibody #1 were introduced into CHO-K1 cells (obtained from Riken Cell Bank). Specifically,  $2 \times 10^5$  CHO-K1 cells were cultured in a Ham's F12 medium (Invitrogen Corp.) containing 1 mL of 10% FBS per well of a 12-well culture plate, and washed with PBS(-). Then, a fresh Ham's F12 medium containing 1 mL of 10% FBS per well was added thereto. 250 ng each of the vectors lysed in 30  $\mu$ L of OptiMEM (Invitrogen Corp.) was mixed with 30  $\mu$ L of Polyfect transfection reagent (Qiagen N.V.), and this mixture was added to each well. The CHO-K1 cells cotransfected with the recombinant vectors were cultured in a Ham's F12 medium containing 10% FBS supplemented with 200 pg/mL Zeocin (Invitrogen Corp.) and 200  $\mu$ g/mL Geneticin (Roche Diagnostics K.K.) and then inoculated to a 96-well plate at a density of 0.5 cells/well to prepare a cell line stably producing a human-mouse chimeric monoclonal antibody #1 (#1) having the variable regions of the mouse monoclonal antibody #1.

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Each prepared cell line was cultured for 5 days in a 150-cm<sup>2</sup> flask at a density of  $5 \times 10^5$  cells/mL using 30 mL of a serum-free OptiCHO medium (Invitrogen Corp.) to obtain culture supernatants containing human-mouse chimeric monoclonal antibody #1.

Also, cell lines stably producing human-mouse chimeric comparative monoclonal antibodies 1 to 11 were prepared in the same way as above using the following anti-CAPRIN-1 mouse-derived monoclonal antibodies described in WO2010/016526 as comparative antibodies: a comparative antibody 1 consisting of the heavy chain variable region of SEQ ID NO: 15 and the light chain variable region of SEQ ID NO: 16; a comparative antibody 2 consisting of the heavy chain variable region of SEQ ID NO: 17 and the light chain variable region of SEQ ID NO: 18; a comparative antibody 3 consisting of the heavy chain variable region of SEQ ID NO: 19 and the light chain variable region of SEQ ID NO: 20; a comparative antibody 4 consisting of the heavy chain variable region of SEQ ID NO: 21 and the light chain variable region of SEQ ID NO: 22; a comparative antibody 5 consisting of the heavy chain variable region of SEQ ID NO: 23 and the light chain variable region of SEQ ID NO: 24; a comparative antibody 6 consisting of the heavy chain variable region of SEQ ID NO: 25 and the light chain variable region of SEQ ID NO: 26; a comparative antibody 7 consisting of the heavy chain variable region of SEQ ID NO: 27 and the light chain variable region of SEQ ID NO: 28; a comparative antibody 8 consisting of the heavy chain variable region of SEQ ID NO: 29 and the light chain variable region of SEQ ID NO: 30; a comparative antibody 9 consisting of the heavy chain variable region of SEQ ID NO: 31 and the light chain variable region of SEQ ID NO: 32; a comparative antibody 10 consisting of the heavy chain variable region of SEQ ID NO: 33 and the light chain variable region of SEQ ID NO: 34; and a comparative antibody 11 consisting of the heavy chain variable region of SEQ ID NO: 35 and the light chain variable region of SEQ ID NO: 36. Each prepared cell line was cultured for 5 days in a 150-cm<sup>2</sup> flask at a density of  $5 \times 10^5$  cells/mL using 30 mL of a serum-free OptiCHO medium (Invitrogen Corp.) to obtain culture supernatants containing any of human-mouse chimeric comparative antibodies 1 to 11.

## Example 5

## Expression of CAPRIN-1 on Surface of Various Cancer Cells Using Anti-CAPRIN-1 Antibody #1

Next, the human breast cancer cell lines (ZR75-1, MCF7, T47D, SK-BR-3, MDA-MB-157, BT-20, MDA-MB-231V, and MRK-nu-1), the kidney cancer cell lines (Caki-1, Caki-2, A498, and ACHN), the urinary bladder cancer cell line (T24), the ovary cancer cell line (SKOV3), the lung cancer cell lines (QG56 and A549), the pancreatic cancer cell lines (Capan-2 and MIAPaCa-2), the prostate cancer cell line (PC3), the uterine cervix cancer cell line (SW756), the fibrosarcoma cell line (HT1080), the brain tumor cell lines (T98G, U87MG, U251, SNB 19, and U373), the gastric cancer cell lines (MNK28 and MNK45), the large bowel cancer cell lines (11T29, Lovo, CaCo2, SW480, and HCT116), the leukemia cell line (AML5), and the lymphoma cell line (Ramos) confirmed to have CAPRIN-1 gene expression were examined for their expression of CAPRIN-1 proteins on the cell surface using the culture supernatants containing #1 obtained in Example 4.  $5 \times 10^5$  cells of each cell line were centrifuged in each 1.5-mL microcentrifuge tube. Each culture supernatant (100  $\mu$ L) containing the antibody #1 was added to the tube and left standing for 1 hour on ice. After washing with PBS,



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HTC-labeled goat anti-mouse IgG (H+L) antibodies (Jackson ImmunoResearch Laboratories, Inc.) diluted with PBS containing 0.1% FBS were added thereto and left standing at 4° C. for 30 minutes. After washing with PBS, the fluorescence intensity was measured using FACS Calibur (Becton, Dickinson and Company). The negative control used was cells reacted only with secondary antibodies. As a result, the cells supplemented with the antibody #1 had fluorescence intensity at least 35% stronger than that of the negative control. This demonstrated that CAPRIN-1 proteins are expressed on the cell membrane surface of the human cancer cell lines. The above rate of enhancement in fluorescence intensity was indicated by the rate of increase in mean fluorescence intensity (MFI) in each cell line and calculated according to the following expression:

$$\text{Rate of increase in mean fluorescence intensity (Rate of enhancement in fluorescence intensity) (\%)} = \frac{(\text{MFI of cells reacted with the anti-CAPRIN-1 antibodies}) - (\text{Control MFI})}{(\text{Control MFI})} \times 100.$$

## Example 6

## Antitumor Effect (ADCC Activity) of Anti-CAPRIN-1 Antibody on Cancer Cell

The anti-CAPRIN-1 human-mouse chimeric monoclonal antibody #1 obtained in Example 4 was studied for its ability to damage CAPRIN-1-expressing cancer cells by ADCC activity assay. The culture supernatant of the cells producing #1 was purified using Hitrap Protein A Sepharose FF (GE Healthcare Bio-Sciences Ltd.). After replacement with PBS (-), the solution was filtered through a 0.22- $\mu\text{m}$  filter (Millipore Corp.). The resulting antibody was used for activity assay.  $10^6$  cells each of the human breast cancer cell line MCF7, the human large bowel cancer cell line HCT-116, the human pancreatic cancer cell line MIAPaCa-2, the human kidney cancer cell line Caki-2, and the human lung cancer cell line QG56 confirmed to have CAPRIN-1 expression were collected into a 50-mL centrifuge tube, to which 100  $\mu\text{Ci}$  of chromium 51 was then added, followed by incubation at 37° C. for 2 hours. Then, the cells were washed three times with an RPMI 1640 medium containing 10% FBS and added at a density of  $2 \times 10^3$  cells/well to each 96-well V-bottom plate to prepare target cells. The purified antibody #1 and the human-mouse chimeric comparative antibodies 1 to 11 obtained in Example 4 were each added thereto at a concentration of 1 pg/well. A cell population containing human NK cells separated using a routine method from human peripheral blood lymphocytes was added to the plate at a density of  $2 \times 10^5$  cells/well and cultured at 37° C. for 4 hours under conditions of 5%  $\text{CO}_2$ . After the culture, the amount of chromium 51 released from damaged tumor cells was measured in the culture supernatant to calculate the cytotoxic activity of each anti-CAPRIN-1 antibody against the cancer cells. The negative control used was cells supplemented with isotype control antibodies. The cell population containing NK cells that was used in this evaluation was prepared as follows: human peripheral blood mononuclear cells separated from human peripheral blood according to a routine method using a specific gravity separation solution Histopaque for human peripheral blood mononuclear cell separation (Sigma-Aldrich Corp.) were reacted with various FITC-labeled antibodies (anti-human CD3 antibody, anti-human CD20 antibody, anti-human CD19 antibody, anti-human CD11c antibody, and anti-HLA-DR antibody (Becton, and Dickinson and Company)), and a cell population unstained with the antibodies was separated using a cell sorter (FACS Vantage SE (Bec-

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ton, and Dickinson and Company)) or human NK cell separation kit (Miltenyi Biotec K.K.). As a result of evaluating cytotoxic activity against the cancer cells, the isotype control antibodies used and the comparative antibodies 1 to 11 used had cytotoxic activity less than 5% against all of the human breast cancer cell line MCF7, the human lung cancer cell line QG56, the human large bowel cancer cell line HCT-116, the human pancreatic cancer cell line MIAPaCa-2, and the human kidney cancer cell line Caki-2. By contrast, the antibody #1 exhibited cytotoxic activity of 21%, 11%, 21%, 27%, and 23% against the human breast cancer cell line MCF7, the human lung cancer cell line QG56, the human large bowel cancer cell line HCT-116, the human pancreatic cancer cell line MIAPaCa-2, and the human kidney cancer cell line Caki-2, respectively. Likewise, the isotype control antibodies used and the comparative antibodies 1 to 11 used had cytotoxic activity less than 4% against all other cancer cells, breast cancer cell lines ZR75-1, T47D, Hs578T, BT-20, SK-BR-3, MDA-MB-231V, and MRK-nu-1, glioma cell lines T98G and U373, a lung cancer cell line A549, kidney cancer cell lines Caki-1 and ACHN, a uterine cervix cancer cell line SW756, a urinary bladder cancer cell line T24, gastric cancer cell lines MKN28 and MKN45, a large bowel cancer cell line SW480, a leukemia cell line AML5, and a lymphoma cell line Ramos. By contrast, the antibody #1 was confirmed to have 10% or higher cytotoxic activity against these cell lines. These results showed that the obtained monoclonal antibody #1 against CAPRIN-1 damages CAPRIN-1-expressing cancer cells through its ADCC activity, and demonstrated that the antibody #1 exhibits stronger cytotoxic activity against human cancer cells than that of the comparative antibodies 1 to 11.

These results about cytotoxic activity were obtained by mixing the antibody against CAPRIN-1 used in the present invention, lymphocyte cells (population containing NK cells), and  $2 \times 10^3$  cells of each cancer cell line with incorporated chromium 51, as described above, followed by culturing the cells for 4 hours, after the culture, measuring the amount of chromium 51 released into the medium, and calculating the cytotoxic activity against each cancer cell line according to the following expression.

$$\text{Expression: Cytotoxic activity (\%)} = \frac{\text{Amount of chromium 51 released from the target cells supplemented with the antibody against CAPRIN-1 and lymphocyte cells (population containing NK cells)}}{\text{Amount of chromium 51 released from target cells supplemented with 1 N hydrochloric acid}} \times 100.$$

## Example 7

## Antitumor Effect of Anti-CAPRIN-1 Monoclonal Antibody on Mouse In Vivo

Next, the human-mouse chimeric monoclonal antibody #1 obtained in Example 4 was evaluated for its antitumor effect on cancer-bearing mice in vivo. The antibody used was column-purified from the culture supernatant of each cell line producing the antibody #1. Similarly, the anti-CAPRIN-1 antibodies human-mouse chimeric comparative monoclonal antibodies 1 to 11 prepared in Example 4 were also evaluated for their antitumor effects on cancer-bearing mice in vivo.

The antibody #1 was studied for its antitumor effect using cancer-bearing mice in which a CAPRIN-1-expressing human-derived cancer cell line was transplanted.  $2 \times 10^6$  human pancreatic cancer cell line Capan-2 cells (purchased from ATCC) per mouse were subcutaneously transplanted into the backs of 65 Balb/c nude mice (Japan SLC, Inc.) and

grown until the size of tumor became approximately 5 mm in diameter. The antibody #1 and the human-mouse chimeric comparative antibodies 1 to 11 were each intraperitoneally administered at a dose of 200  $\mu$ g (200  $\mu$ l/mouse to 5 (per antibody) of these cancer-bearing mice. Then, each antibody was intraperitoneally administered to the cancer-bearing mice at the same dose as above a total of three times for 2 days. The size of tumor was measured every day, and the antitumor effect was observed. On the other hand, PBS(-) was administered instead of the antibodies to the remaining 5 cancer-bearing mice, which were in turn used as a control group. The size of tumor was calculated in terms of volume according to the expression  $0.5 \times (\text{Major axis} \times \text{Minor axis} \times \text{Minor axis})$ .

As a result of observing the antitumor effect, in the test group that received the antibody #1 against CAPRIN-1, tumor proliferation was reduced to 65% at day 29 after the antibody administration (with the tumor size in the control group at the same date defined as 100%). By contrast, in the mice that received the human-mouse chimeric comparative antibodies 1 to 11, tumor proliferation was reduced to approximately 85%. These results demonstrated that the obtained antibody #1 against CAPRIN-1 exerts an in vivo antitumor effect on CAPRIN-1-expressing cancer cells. These results also demonstrated that the antibody #1 exerts a stronger in vivo antitumor effect than that of the comparative antibodies 1 to 11.

#### Example 8

##### The Number of CAPRIN-1 Molecules on Surface of Various Cancer Cells Recognized by Anti-CAPRIN-1 Antibody #1

Human breast cancer cell lines (ZR75-1, MCF7, T47D, SK-BR-3, MDA-MB-157, BT-20, MDA-MB-231V, and MRK-nu-1), kidney cancer cell lines (Caki-1, Caki-2, A498, and ACHN), a urinary bladder cancer cell line (T24), an ovary cancer cell line (SKOV3), lung cancer cell lines (QG56 and A549), pancreatic cancer cell lines (MIAPaCa-2 and Capan-2), a prostate cancer cell line (PC3), a uterine cervix cancer

cell line (SW756), a fibrosarcoma cell line (HT1080), brain tumor cell lines (T98G, U87MG, U251, SNB19, and U373), gastric cancer cell lines (MNK28 and MNK45), large bowel cancer cell lines (HT29, Lovo, CaCo2, SW480, and HCT116), a leukemia cell line (AML5), and a lymphoma cell line (Ramos) were examined using an assay kit QIFIKIT for the number of molecules (Dako Japan Inc.) for the number of CAPRIN-1 molecules on their cell surface recognized by the anti-CAPRIN-1 antibody #1. Similarly, the number of CAPRIN-1 molecules on the surface of these various cancer cells was also examined using the comparative antibodies 1 to 11, which are anti-CAPRIN-1 monoclonal antibodies prepared in Example 4.

According to the protocol attached to the kit, the antibody #1 and comparative antibodies 1 to 11 were diluted into 5  $\mu$ g/mL (in terms of final concentration) with PBS, and this dilution was added to each cell line and reacted for 30 minutes. After washing with PBS, fluorescently labeled anti-mouse IgG antibodies attached to the kit were added as secondary antibodies, together with calibration beads attached to the kit, to each cell line and left standing for 45 minutes on ice. Each cell line and the calibration beads were washed with PBS. Then, the fluorescence intensity was measured using FACS Calibur (Becton, Dickinson and Company) to obtain a mean fluorescence intensity value (mean). Further, comparative antibodies are measured similarly as above to obtain a mean. The negative control used was cells reacted with isotype control antibodies, and a mean was also obtained. Each mean fluorescence intensity value (mean) was used to calculate the number of molecules according to the protocol attached to the kit. As a result, the number of CAPRIN-1 molecules on the surface of various cancer cells recognized by the antibody #1 was  $10^5$  or more per cell for all the examined human cancer cell lines. On the other hand, the number of molecules recognized by the comparative monoclonal antibodies 1 to 11 was less than  $10^5$  per cell.

##### Industrial Applicability

The antibody of the present invention is useful in the treatment and/or prevention of cancer.

All publications, patents, and patent applications cited herein are incorporated herein by reference in their entirety.

#### SEQUENCE LISTING

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<210> SEQ ID NO 1

<211> LENGTH: 5562

<212> TYPE: DNA

<213> ORGANISM: Homo sapiens

<220> FEATURE:

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<222> LOCATION: (190)..(2319)

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1 5 10

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gct ggg aca agc aaa cct tta cat agc agt gga atc aat gta aat gca Ala Gly Thr Ser Lys Pro Leu His Ser Ser Gly Ile Asn Val Asn Ala 495 500 505 510				1719
gct cca ttc caa tcc atg caa acg gtg ttc aat atg aat gcc cca gtt Ala Pro Phe Gln Ser Met Gln Thr Val Phe Asn Met Asn Ala Pro Val 515 520 525				1767
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cct cag cag aac act gga ttt cca cgt agc aat cag ccc tat tac aat Pro Gln Gln Asn Thr Gly Phe Pro Arg Ser Asn Gln Pro Tyr Tyr Asn 595 600 605				2007
agt cgt ggt gtg tct cgt gga ggc tcc cgt ggt gct aga ggc ttg atg Ser Arg Gly Val Ser Arg Gly Gly Ser Arg Gly Ala Arg Gly Leu Met 610 615 620				2055
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tac cgc cct tca ttc tct aac act cca aac agt ggt tat aca cag tct Tyr Arg Pro Ser Phe Ser Asn Thr Pro Asn Ser Gly Tyr Thr Gln Ser 640 645 650				2151
cag ttc agt gct ccc cgg gat tac tct ggc tat caa cgg gat gga tat				2199

Gln 655	Phe	Ser	Ala	Pro	Arg 660	Asp	Tyr	Ser	Gly	Tyr 665	Gln	Arg	Asp	Gly	Tyr 670	
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cca Pro	cga Arg	ggt Gly	cgt Arg 690	gga Gly	ggg Gly	ccc Pro	cca Pro	aga Gly 695	ccc Pro	aac Asn	aga Arg	ggg Gly	atg Met 700	ccg Pro	caa Gln	2295
atg Met 705	aac Asn	act Thr	cag Gln	caa Gln	gtg Val	aat Asn	taa	tctgattcac	aggattatgt	ttaatcgcca						2349
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&lt;210&gt; SEQ ID NO 2

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&lt;212&gt; TYPE: PRT

&lt;213&gt; ORGANISM: Homo sapiens

&lt;400&gt; SEQUENCE: 2

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Gly Ala Ala Ala Pro Ala Ser Gln His Pro Ala Thr Gly Thr Gly Ala
35          40          45
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50          55          60
Leu Arg Asn Leu Glu Lys Lys Lys Gly Lys Leu Asp Asp Tyr Gln Glu
65          70          75          80
Arg Met Asn Lys Gly Glu Arg Leu Asn Gln Asp Gln Leu Asp Ala Val
85          90          95
Ser Lys Tyr Gln Glu Val Thr Asn Asn Leu Glu Phe Ala Lys Glu Leu
100         105         110
Gln Arg Ser Phe Met Ala Leu Ser Gln Asp Ile Gln Lys Thr Ile Lys
115         120         125
Lys Thr Ala Arg Arg Glu Gln Leu Met Arg Glu Glu Ala Glu Gln Lys
130         135         140

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Asp	Asp	Glu	Val	Arg	Thr	Asp	Leu	Lys	Gln	Gly	Leu	Asn	Gly	Val	Pro	165	170	175	
Ile	Leu	Ser	Glu	Glu	Glu	Leu	Ser	Leu	Leu	Asp	Glu	Phe	Tyr	Lys	Leu	180	185	190	
Val	Asp	Pro	Glu	Arg	Asp	Met	Ser	Leu	Arg	Leu	Asn	Glu	Gln	Tyr	Glu	195	200	205	
His	Ala	Ser	Ile	His	Leu	Trp	Asp	Leu	Leu	Glu	Gly	Lys	Glu	Lys	Pro	210	215	220	
Val	Cys	Gly	Thr	Thr	Tyr	Lys	Val	Leu	Lys	Glu	Ile	Val	Glu	Arg	Val	225	230	235	240
Phe	Gln	Ser	Asn	Tyr	Phe	Asp	Ser	Thr	His	Asn	His	Gln	Asn	Gly	Leu	245	250	255	
Cys	Glu	Glu	Glu	Glu	Ala	Ala	Ser	Ala	Pro	Ala	Val	Glu	Asp	Gln	Val	260	265	270	
Pro	Glu	Ala	Glu	Pro	Glu	Pro	Ala	Glu	Glu	Tyr	Thr	Glu	Gln	Ser	Glu	275	280	285	
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Ser	Val	Pro	Glu	Pro	His	Ser	Leu	Thr	Pro	Val	Ala	Gln	Ala	Asp	Pro	340	345	350	
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Pro	Tyr	Asn	Phe	Ile	Gln	Asp	Ser	Met	Leu	Asp	Phe	Glu	Asn	Gln	Thr	370	375	380	
Leu	Asp	Pro	Ala	Ile	Val	Ser	Ala	Gln	Pro	Met	Asn	Pro	Thr	Gln	Asn	385	390	395	400
Met	Asp	Met	Pro	Gln	Leu	Val	Cys	Pro	Pro	Val	His	Ser	Glu	Ser	Arg	405	410	415	
Leu	Ala	Gln	Pro	Asn	Gln	Val	Pro	Val	Gln	Pro	Glu	Ala	Thr	Gln	Val	420	425	430	
Pro	Leu	Val	Ser	Ser	Thr	Ser	Glu	Gly	Tyr	Thr	Ala	Ser	Gln	Pro	Leu	435	440	445	
Tyr	Gln	Pro	Ser	His	Ala	Thr	Glu	Gln	Arg	Pro	Gln	Lys	Glu	Pro	Ile	450	455	460	
Asp	Gln	Ile	Gln	Ala	Thr	Ile	Ser	Leu	Asn	Thr	Asp	Gln	Thr	Thr	Ala	465	470	475	480
Ser	Ser	Ser	Leu	Pro	Ala	Ala	Ser	Gln	Pro	Gln	Val	Phe	Gln	Ala	Gly	485	490	495	
Thr	Ser	Lys	Pro	Leu	His	Ser	Ser	Gly	Ile	Asn	Val	Asn	Ala	Ala	Pro	500	505	510	
Phe	Gln	Ser	Met	Gln	Thr	Val	Phe	Asn	Met	Asn	Ala	Pro	Val	Pro	Pro	515	520	525	
Val	Asn	Glu	Pro	Glu	Thr	Leu	Lys	Gln	Gln	Asn	Gln	Tyr	Gln	Ala	Ser	530	535	540	
Tyr	Asn	Gln	Ser	Phe	Ser	Ser	Gln	Pro	His	Gln	Val	Glu	Gln	Thr	Glu	545	550	555	560
Leu	Gln	Gln	Glu	Gln	Leu	Gln	Thr	Val	Val	Gly	Thr	Tyr	His	Gly	Ser				

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Gln Asn Thr Gly Phe Pro Arg Ser Asn Gln Pro Tyr Tyr Asn Ser Arg 595 600 605		
Gly Val Ser Arg Gly Gly Ser Arg Gly Ala Arg Gly Leu Met Asn Gly 610 615 620		
Tyr Arg Gly Pro Ala Asn Gly Phe Arg Gly Gly Tyr Asp Gly Tyr Arg 625 630 635 640		
Pro Ser Phe Ser Asn Thr Pro Asn Ser Gly Tyr Thr Gln Ser Gln Phe 645 650 655		
Ser Ala Pro Arg Asp Tyr Ser Gly Tyr Gln Arg Asp Gly Tyr Gln Gln 660 665 670		
Asn Phe Lys Arg Gly Ser Gly Gln Ser Gly Pro Arg Gly Ala Pro Arg 675 680 685		
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ggaagggacc gccacccttg cccctcagc tgcccactcg tgatttccag cggcctccgc	180	
gcgcgcacg atg ccc tcg gcc acc agc cac agc ggg agc ggc agc aag tcg Met Pro Ser Ala Thr Ser His Ser Gly Ser Gly Ser Lys Ser 1 5 10	231	
tcc gga ccg cca ccg ccg tcg ggt tcc tcc ggg agt gag gcg gcc gcg Ser Gly Pro Pro Pro Pro Ser Gly Ser Ser Gly Ser Glu Ala Ala Ala 15 20 25 30	279	
gga gcc ggg gcc gcc gcg ccg gct tct cag cac ccc gca acc ggc acc Gly Ala Gly Ala Ala Ala Pro Ala Ser Gln His Pro Ala Thr Gly Thr 35 40 45	327	
ggc gct gtc cag acc gag gcc atg aag cag att ctc ggg gtg atc gac Gly Ala Val Gln Thr Glu Ala Met Lys Gln Ile Leu Gly Val Ile Asp 50 55 60	375	
aag aaa ctt ccg aac ctg gag aag aaa aag ggt aag ctt gat gat tac Lys Lys Leu Arg Asn Leu Glu Lys Lys Lys Gly Lys Leu Asp Asp Tyr 65 70 75	423	
cag gaa cga atg aac aaa ggg gaa agg ctt aat caa gat cag ctg gat Gln Glu Arg Met Asn Lys Gly Glu Arg Leu Asn Gln Asp Gln Leu Asp 80 85 90	471	
gcc gtt tct aag tac cag gaa gtc aca aat aat ttg gag ttt gca aaa Ala Val Ser Lys Tyr Gln Glu Val Thr Asn Asn Leu Glu Phe Ala Lys 95 100 105 110	519	
gaa tta cag agg agt ttc atg gca cta agt caa gat att cag aaa aca Glu Leu Gln Arg Ser Phe Met Ala Leu Ser Gln Asp Ile Gln Lys Thr 115 120 125	567	



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Ile Lys Lys Thr Ala Arg Arg Glu Gln Leu Met Arg Glu Glu Ala Glu	
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Gln Lys Arg Leu Lys Thr Val Leu Glu Leu Gln Tyr Val Leu Asp Lys	
145 150 155	
ttg gga gat gat gaa gtg cgg act gac ctg aaa caa ggt ttg aat gga	711
Leu Gly Asp Asp Glu Val Arg Thr Asp Leu Lys Gln Gly Leu Asn Gly	
160 165 170	
gtg cca ata ttg tcc gaa gag gag ttg tca ttg ttg gat gaa ttc tat	759
Val Pro Ile Leu Ser Glu Glu Glu Leu Ser Leu Leu Asp Glu Phe Tyr	
175 180 185 190	
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Lys Leu Val Asp Pro Glu Arg Asp Met Ser Leu Arg Leu Asn Glu Gln	
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Tyr Glu His Ala Ser Ile His Leu Trp Asp Leu Leu Glu Gly Lys Glu	
210 215 220	
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Lys Pro Val Cys Gly Thr Thr Tyr Lys Val Leu Lys Glu Ile Val Glu	
225 230 235	
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Arg Val Phe Gln Ser Asn Tyr Phe Asp Ser Thr His Asn His Gln Asn	
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Gly Leu Cys Glu Glu Glu Ala Ala Ser Ala Pro Ala Val Glu Asp	
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Gln Val Pro Glu Ala Glu Pro Glu Pro Ala Glu Glu Tyr Thr Glu Gln	
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Glu Thr Val Glu Val Val Asn Ser Leu Gln Gln Gln Pro Gln Ala Ala	
320 325 330	
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Ser Pro Ser Val Pro Glu Pro His Ser Leu Thr Pro Val Ala Gln Ala	
335 340 345 350	
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Asp Pro Leu Val Arg Arg Gln Arg Val Gln Asp Leu Met Ala Gln Met	
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370 375 380	
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Gln Thr Leu Asp Pro Ala Ile Val Ser Ala Gln Pro Met Asn Pro Thr	
385 390 395	
caa aac atg gac atg ccc cag ctg gtt tgc cct cca gtt cat tct gaa	1431
Gln Asn Met Asp Met Pro Gln Leu Val Cys Pro Pro Val His Ser Glu	
400 405 410	
tct aga ctt gct cag cct aat caa gtt cct gta caa cca gaa gcg aca	1479
Ser Arg Leu Ala Gln Pro Asn Gln Val Pro Val Gln Pro Glu Ala Thr	
415 420 425 430	
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Gln Val Pro Leu Val Ser Ser Thr Ser Glu Gly Tyr Thr Ala Ser Gln	
435 440 445	

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Pro Ile Asp Gln Ile Gln Ala Thr Ile Ser Leu Asn Thr Asp Gln Thr	
465 470 475	
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Thr Ala Ser Ser Ser Leu Pro Ala Ala Ser Gln Pro Gln Val Phe Gln	
480 485 490	
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Ala Gly Thr Ser Lys Pro Leu His Ser Ser Gly Ile Asn Val Asn Ala	
495 500 505 510	
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Pro Pro Val Asn Glu Pro Glu Thr Leu Lys Gln Gln Asn Gln Tyr Gln	
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Thr Glu Leu Gln Gln Glu Gln Leu Gln Thr Val Val Gly Thr Tyr His	
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Gly Ser Pro Asp Gln Ser His Gln Val Thr Gly Asn His Gln Gln Pro	
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610 615 620	
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Asn Gly Tyr Arg Gly Pro Ala Asn Gly Phe Arg Gly Gly Tyr Asp Gly	
625 630 635	
tac cgc cct tca ttc tct aac act cca aac agt ggt tat aca cag tct	2151
Tyr Arg Pro Ser Phe Ser Asn Thr Pro Asn Ser Gly Tyr Thr Gln Ser	
640 645 650	
cag ttc agt gct ccc cgg gat tac tct ggc tat caa cgg gat gga tat	2199
Gln Phe Ser Ala Pro Arg Asp Tyr Ser Gly Tyr Gln Arg Asp Gly Tyr	
655 660 665 670	
cag cag aat ttc aag cga ggc tct ggg cag agt gga cca cgg gga gcc	2247
Gln Gln Asn Phe Lys Arg Gly Ser Gly Gln Ser Gly Pro Arg Gly Ala	
675 680 685	
cca cga ggt aat att ttg tgg tgg tga tctagctcc taagtggagc	2294
Pro Arg Gly Asn Ile Leu Trp Trp	
690	
ttctgttctg gccttggaag agctgttaat agtctgcatg ttaggaatac atttatectt	2354
tccagacttg ttgctagga ttaaatgaaa tgctctgttt ctaaaactta atcttggaac	2414
caaatTTTaa tttttgaatg atttaatttt cctgtttact atataaactg tcttgaaaac	2474
tagaacatat tctcttctca gaaaaagtgt ttttccaact gaaaattatt tttcagggcc	2534
taaaacctgc taaatgtttt taggaagtac ttactgaaac atttttgtaa gacatttttg	2594
gaatgagatt gaacatttat ataaatttat tattcctctt tcattttttt gaaacatgcc	2654
tattatattt tagggccaga caccctttta tggccggata agccatagtt aacattttaga	2714

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gaaccattta gaagtgatag aactaatgga atttgcaatg ccttttggac ctctattagt 2774
gatataaata tcaagttatt tctgactttt aaacaaaact cccaaattcc taacttattg 2834
agctatactt aaaaaaaatt acaggtttag agagtgtttt gtttttcttt tactgttgga 2894
aaactacttc ccattttggc aggaagttaa cctatttaac aattagagct agcatttcac 2954
gtagtctgaa attctaaatg gttctctgat ttgaggagg ttaaacatca aacagggttc 3014
ctctattggc cataacatgt ataaaaatg tgtaaggag gaattacaac gtactttgat 3074
ttgaatacta gtagaaactg gccaggaaaa aggtacattt ttctaaaaat taatggatca 3134
cttggaatt actgacttga ctagaagtat caaaggatgt ttgcatgtga atgtgggtta 3194
tggtctttcc cacctgttag catattgat gaaagttgag ttaactgata gctaaaaatc 3254
tgttttaaca gcatgtaaaa agttatttta tctgttaaaa gtcattatac agttttgaat 3314
gttatgtagt ttctttttaa cagtttaggt aataaggtct gttttcattc tgggtgctttt 3374
attaattttg atagtatgat gttacttact actgaaatgt aagctagagt gtacactaga 3434
atgtaagctc catgagagca ggtaccttgt ctgtcttctc tgctgtatct attccaacg 3494
cttgatgatg gtgcctggca catagtaggc actcaataaa tatttgttga atgaatgaa 3553

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<210> SEQ ID NO 4
<211> LENGTH: 694
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens

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<400> SEQUENCE: 4

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```

Met Pro Ser Ala Thr Ser His Ser Gly Ser Gly Ser Lys Ser Ser Gly
1          5          10          15
Pro Pro Pro Pro Ser Gly Ser Ser Gly Ser Glu Ala Ala Ala Gly Ala
20         25         30
Gly Ala Ala Ala Pro Ala Ser Gln His Pro Ala Thr Gly Thr Gly Ala
35         40         45
Val Gln Thr Glu Ala Met Lys Gln Ile Leu Gly Val Ile Asp Lys Lys
50         55         60
Leu Arg Asn Leu Glu Lys Lys Lys Gly Lys Leu Asp Asp Tyr Gln Glu
65         70         75         80
Arg Met Asn Lys Gly Glu Arg Leu Asn Gln Asp Gln Leu Asp Ala Val
85         90         95
Ser Lys Tyr Gln Glu Val Thr Asn Asn Leu Glu Phe Ala Lys Glu Leu
100        105        110
Gln Arg Ser Phe Met Ala Leu Ser Gln Asp Ile Gln Lys Thr Ile Lys
115        120        125
Lys Thr Ala Arg Arg Glu Gln Leu Met Arg Glu Glu Ala Glu Gln Lys
130        135        140
Arg Leu Lys Thr Val Leu Glu Leu Gln Tyr Val Leu Asp Lys Leu Gly
145        150        155        160
Asp Asp Glu Val Arg Thr Asp Leu Lys Gln Gly Leu Asn Gly Val Pro
165        170        175
Ile Leu Ser Glu Glu Glu Leu Ser Leu Leu Asp Glu Phe Tyr Lys Leu
180        185        190
Val Asp Pro Glu Arg Asp Met Ser Leu Arg Leu Asn Glu Gln Tyr Glu
195        200        205
His Ala Ser Ile His Leu Trp Asp Leu Leu Glu Gly Lys Glu Lys Pro
210        215        220
Val Cys Gly Thr Thr Tyr Lys Val Leu Lys Glu Ile Val Glu Arg Val

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225					230						235					240
Phe	Gln	Ser	Asn	Tyr	Phe	Asp	Ser	Thr	His	Asn	His	Gln	Asn	Gly	Leu	
				245					250					255		
Cys	Glu	Glu	Glu	Glu	Ala	Ala	Ser	Ala	Pro	Ala	Val	Glu	Asp	Gln	Val	
			260					265					270			
Pro	Glu	Ala	Glu	Pro	Glu	Pro	Ala	Glu	Glu	Tyr	Thr	Glu	Gln	Ser	Glu	
		275					280					285				
Val	Glu	Ser	Thr	Glu	Tyr	Val	Asn	Arg	Gln	Phe	Met	Ala	Glu	Thr	Gln	
	290					295					300					
Phe	Thr	Ser	Gly	Glu	Lys	Glu	Gln	Val	Asp	Glu	Trp	Thr	Val	Glu	Thr	
305					310					315					320	
Val	Glu	Val	Val	Asn	Ser	Leu	Gln	Gln	Gln	Pro	Gln	Ala	Ala	Ser	Pro	
				325					330					335		
Ser	Val	Pro	Glu	Pro	His	Ser	Leu	Thr	Pro	Val	Ala	Gln	Ala	Asp	Pro	
		340						345						350		
Leu	Val	Arg	Arg	Gln	Arg	Val	Gln	Asp	Leu	Met	Ala	Gln	Met	Gln	Gly	
		355					360					365				
Pro	Tyr	Asn	Phe	Ile	Gln	Asp	Ser	Met	Leu	Asp	Phe	Glu	Asn	Gln	Thr	
	370					375					380					
Leu	Asp	Pro	Ala	Ile	Val	Ser	Ala	Gln	Pro	Met	Asn	Pro	Thr	Gln	Asn	
385					390					395					400	
Met	Asp	Met	Pro	Gln	Leu	Val	Cys	Pro	Pro	Val	His	Ser	Glu	Ser	Arg	
				405					410					415		
Leu	Ala	Gln	Pro	Asn	Gln	Val	Pro	Val	Gln	Pro	Glu	Ala	Thr	Gln	Val	
			420					425					430			
Pro	Leu	Val	Ser	Ser	Thr	Ser	Glu	Gly	Tyr	Thr	Ala	Ser	Gln	Pro	Leu	
	435						440					445				
Tyr	Gln	Pro	Ser	His	Ala	Thr	Glu	Gln	Arg	Pro	Gln	Lys	Glu	Pro	Ile	
	450					455					460					
Asp	Gln	Ile	Gln	Ala	Thr	Ile	Ser	Leu	Asn	Thr	Asp	Gln	Thr	Thr	Ala	
465					470					475					480	
Ser	Ser	Ser	Leu	Pro	Ala	Ala	Ser	Gln	Pro	Gln	Val	Phe	Gln	Ala	Gly	
				485					490					495		
Thr	Ser	Lys	Pro	Leu	His	Ser	Ser	Gly	Ile	Asn	Val	Asn	Ala	Ala	Pro	
		500						505					510			
Phe	Gln	Ser	Met	Gln	Thr	Val	Phe	Asn	Met	Asn	Ala	Pro	Val	Pro	Pro	
	515							520				525				
Val	Asn	Glu	Pro	Glu	Thr	Leu	Lys	Gln	Gln	Asn	Gln	Tyr	Gln	Ala	Ser	
	530					535						540				
Tyr	Asn	Gln	Ser	Phe	Ser	Ser	Gln	Pro	His	Gln	Val	Glu	Gln	Thr	Glu	
545					550					555					560	
Leu	Gln	Gln	Glu	Gln	Leu	Gln	Thr	Val	Val	Gly	Thr	Tyr	His	Gly	Ser	
				565					570					575		
Pro	Asp	Gln	Ser	His	Gln	Val	Thr	Gly	Asn	His	Gln	Gln	Pro	Pro	Gln	
		580						585					590			
Gln	Asn	Thr	Gly	Phe	Pro	Arg	Ser	Asn	Gln	Pro	Tyr	Tyr	Asn	Ser	Arg	
		595					600					605				
Gly	Val	Ser	Arg	Gly	Gly	Ser	Arg	Gly	Ala	Arg	Gly	Leu	Met	Asn	Gly	
	610					615					620					
Tyr	Arg	Gly	Pro	Ala	Asn	Gly	Phe	Arg	Gly	Gly	Tyr	Asp	Gly	Tyr	Arg	
625					630					635					640	
Pro	Ser	Phe	Ser	Asn	Thr	Pro	Asn	Ser	Gly	Tyr	Thr	Gln	Ser	Gln	Phe	
				645					650					655		

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Ser Ala Pro Arg Asp Tyr Ser Gly Tyr Gln Arg Asp Gly Tyr Gln Gln  
                   660                                  665                                  670

Asn Phe Lys Arg Gly Ser Gly Gln Ser Gly Pro Arg Gly Ala Pro Arg  
                   675                                  680                                  685

Gly Asn Ile Leu Trp Trp  
                   690

<210> SEQ ID NO 5  
 <211> LENGTH: 5  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 5

Asn Ser Trp Phe Asn  
 1                                  5

<210> SEQ ID NO 6  
 <211> LENGTH: 19  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 6

Glu Ile Arg Leu Thr Ser Asp Asn Tyr Ala Ile Tyr Tyr Ala Glu Ser  
 1                                  5                                  10                                  15

Val Lys Gly

<210> SEQ ID NO 7  
 <211> LENGTH: 11  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 7

Pro Glu Thr Ala Arg Ala Thr Phe Ala Tyr Trp  
 1                                  5                                  10

<210> SEQ ID NO 8  
 <211> LENGTH: 114  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 8

Gly Gly Gly Leu Val Gln Pro Gly Gly Ser Met Arg Leu Ser Cys Val  
 1                                  5                                  10                                  15

Ala Ser Gly Phe Thr Phe Ser Asn Ser Trp Phe Asn Trp Val Arg Gln  
                   20                                  25                                  30

Ser Pro Glu Lys Gly Leu Glu Trp Val Ala Glu Ile Arg Leu Thr Ser  
                   35                                  40                                  45

Asp Asn Tyr Ala Ile Tyr Tyr Ala Glu Ser Val Lys Gly Arg Phe Thr  
                   50                                  55                                  60

Ile Ser Arg Asp Asp Ser Lys Ser Ser Val Tyr Leu Gln Met Asn Asn  
                   65                                  70                                  75                                  80

Leu Arg Ala Glu Asp Thr Gly Ile Tyr Tyr Cys Thr Arg Pro Glu Thr  
                   85                                  90                                  95

Ala Arg Ala Thr Phe Ala Tyr Trp Gly Gln Gly Thr Thr Val Thr Val  
                   100                                  105                                  110

Ser Ser

<210> SEQ ID NO 9  
 <211> LENGTH: 15

-continued

<212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 9

Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Ile Val His Ser Asn  
 1 5 10 15

<210> SEQ ID NO 10  
 <211> LENGTH: 7  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 10

Lys Val Phe Asn Arg Phe Ser  
 1 5

<210> SEQ ID NO 11  
 <211> LENGTH: 9  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 11

Phe Gln Gly Ser His Val Pro Arg Thr  
 1 5

<210> SEQ ID NO 12  
 <211> LENGTH: 118  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 12

Pro Ala Ser Thr Ser Asp Val Leu Met Thr Gln Thr Pro Leu Ser Leu  
 1 5 10 15

Pro Val Ser Leu Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln  
 20 25 30

Ser Ile Val His Ser Asn Gly Asn Thr Tyr Leu Glu Trp Tyr Leu Gln  
 35 40 45

Lys Pro Gly Gln Ser Pro Lys Val Leu Ile Tyr Lys Val Phe Asn Arg  
 50 55 60

Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp  
 65 70 75 80

Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Leu Gly Val Tyr  
 85 90 95

Tyr Cys Phe Gln Gly Ser His Val Pro Arg Thr Phe Gly Gly Gly Thr  
 100 105 110

Lys Leu Asn Gln Thr Gly  
 115

<210> SEQ ID NO 13  
 <211> LENGTH: 342  
 <212> TYPE: DNA  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 13

ggaggaggct tgggtgcaacc tggaggatct atgagactct cctgtgttgc ctctggattc 60

acttttcagta actcctgggt taactgggtc cgccagtctc cagagaaggg gcttgagtgg 120

gttgctgaaa ttagattgac atctgataat tatgcaatat attatgcgga gtctgtgaaa 180

gggaggttca ccatctcaag agatgattcc aaaagcagtg tctatctgca aatgaacaac 240

ttaagagctg aagacactgg catttattac tgtaccaggc ctgagacagc tcgggctacg 300

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tttgcttact ggggccaagg gaccacggtc acggtctcct ca 342

<210> SEQ ID NO 14  
 <211> LENGTH: 354  
 <212> TYPE: DNA  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 14

```
cctgcttcca ccagtgatgt ttgatgacc caaactccac tctccctgcc tgtcagtctt    60
ggagatcaag cctccatctc ttgcagatct agtcagagca ttgtacatag taatggaaac    120
acctatttag aatggtaact gcagaaacca ggccagtctc caaaggctctt gatctacaaa    180
gtttttaacc gattttctgg ggtcccagac aggttcagtg gcagtggatc agggacagat    240
ttcacactca agatcagcag agtggaggct gaggatctgg gagtttatta ctgctttcaa    300
ggttcacatg ttcctcggac gttcgggtgga ggcaccaagc tgaatcagac gggc      354
```

<210> SEQ ID NO 15  
 <211> LENGTH: 148  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 15

```
Met Glu Trp Ser Gly Val Phe Ile Phe Leu Leu Ser Gly Thr Ala Gly
 1             5             10             15
Val Leu Ser Glu Val Gln Leu His Gln Phe Gly Ala Glu Leu Val Lys
 20            25            30
Pro Gly Ala Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Thr Phe
 35            40            45
Thr Asp Tyr Asn Met Asp Trp Val Lys Gln Ser His Gly Lys Ser Leu
 50            55            60
Glu Trp Ile Gly Asp Ile Asn Pro Asn Tyr Asp Ser Thr Ser Tyr Asn
 65            70            75            80
Gln Lys Phe Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser
 85            90            95
Thr Ala Tyr Met Glu Leu Arg Ser Leu Thr Ser Glu Asp Thr Ala Val
100           105           110
Tyr Tyr Cys Ala Arg Ser Arg Ser Tyr Asp Tyr Glu Gly Phe Ala Tyr
115           120           125
Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ala Ala Lys Thr Thr Pro
130           135           140
Pro Ser Val Tyr
145
```

<210> SEQ ID NO 16  
 <211> LENGTH: 139  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 16

```
Met Ser Val Leu Thr Gln Val Leu Gly Leu Leu Leu Trp Leu Thr
 1             5             10             15
Gly Ala Arg Cys Asp Ile Gln Met Thr Gln Ser Pro Ala Ser Leu Ser
 20            25            30
Ala Ser Val Gly Glu Thr Val Thr Ile Thr Cys Arg Ala Ser Gly Asn
 35            40            45
Ile His Asn Tyr Leu Ala Trp Tyr Gln Gln Lys Gln Gly Lys Ser Pro
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50	55	60
Gln Leu Leu Val Tyr Asn Ala Lys Thr Leu Ala Asp Gly Val Pro Ser		
65	70	75 80
Arg Phe Ser Gly Ser Gly Ser Gly Thr Gln Tyr Ser Leu Lys Ile Asn		
	85	90 95
Ser Leu Gln Pro Glu Asp Phe Gly Ser Tyr Tyr Cys Gln His Phe Trp		
	100	105 110
Ser Thr Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys Arg Ala		
	115	120 125
Asp Ala Ala Pro Thr Val Ser Asn Pro Tyr Asp		
130	135	

<210> SEQ ID NO 17  
 <211> LENGTH: 148  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 17

Met Glu Trp Ser Gly Val Phe Ile Phe Leu Leu Ser Gly Thr Ala Gly
1 5 10 15
Val Leu Ser Glu Val Gln Leu His Gln Phe Gly Ala Glu Leu Val Lys
20 25 30
Pro Gly Ala Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Thr Phe
35 40 45
Thr Asp Tyr Asn Met Asp Trp Val Lys Gln Ser His Gly Lys Ser Leu
50 55 60
Glu Trp Ile Gly Asp Ile Asn Pro Asn Tyr Asp Ser Thr Ser Tyr Asn
65 70 75 80
Gln Lys Phe Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser
85 90 95
Thr Ala Tyr Met Glu Leu Arg Ser Leu Thr Ser Glu Asp Thr Ala Val
100 105 110
Tyr Tyr Cys Ala Arg Ser Arg Ser Tyr Asp Tyr Glu Gly Phe Ala Tyr
115 120 125
Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ala Ala Lys Thr Thr Pro
130 135 140
Pro Ser Val Tyr
145

<210> SEQ ID NO 18  
 <211> LENGTH: 132  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 18

Ala Val Leu Arg Cys Ser Arg Gly Leu Leu Val Ile Trp Ile Ser Asp
1 5 10 15
Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ala Val Thr Ala Gly Glu
20 25 30
Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu Leu Trp Ser Val
35 40 45
Asn Gln Lys Asn Tyr Leu Ser Trp Tyr Gln Gln Lys Gln Arg Gln Pro
50 55 60
Pro Lys Leu Leu Ile Tyr Gly Ala Ser Ile Arg Glu Ser Trp Val Pro
65 70 75 80
Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile



85								90					95				
Ser	Asn	Val	His	Ala	Glu	Asp	Leu	Ala	Val	Tyr	Tyr	Cys	Gln	His	Asn		
100							105					110					
His	Gly	Ser	Phe	Leu	Pro	Ser	Arg	Ser	Glu	Gln	Val	Pro	Ser	Trp	Arg		
115							120					125					
Ser	Asn	Asn	Arg														
130																	

```
<210> SEQ ID NO 19
<211> LENGTH: 148
<212> TYPE: PRT
<213> ORGANISM: Mus musculus
```

<400> SEQUENCE: 19

[illegible]

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<210> SEQ ID NO 20
<211> LENGTH: 117
<212> TYPE: PRT
<213> ORGANISM: Mus musculus
```

&lt;400&gt; SEQUENCE: 20

[illegible]

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115

<210> SEQ ID NO 21  
 <211> LENGTH: 148  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 21

```
Met Glu Trp Ser Gly Val Phe Ile Phe Leu Leu Ser Gly Thr Ala Gly
 1             5             10             15
Val Leu Ser Glu Val Gln Leu His Gln Phe Gly Ala Glu Leu Val Lys
          20             25             30
Pro Gly Ala Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Thr Phe
          35             40             45
Thr Asp Tyr Asn Met Asp Trp Val Lys Gln Ser His Gly Lys Ser Leu
          50             55             60
Glu Trp Ile Gly Asp Ile Asn Pro Asn Tyr Asp Ser Thr Ser Tyr Asn
 65             70             75             80
Gln Lys Phe Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser
          85             90             95
Thr Ala Tyr Met Glu Leu Arg Ser Leu Thr Ser Glu Asp Thr Ala Val
          100            105            110
Tyr Tyr Cys Ala Arg Ser Arg Ser Tyr Asp Tyr Glu Gly Phe Ala Tyr
          115            120            125
Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ala Ala Lys Thr Thr Pro
          130            135            140
Pro Ser Val Tyr
145
```

<210> SEQ ID NO 22  
 <211> LENGTH: 94  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 22

```
Ser Gly Asp Arg Val Ser Leu Ser Cys Arg Ala Ser Gln Ser Ile Ser
 1             5             10             15
Asn Tyr Leu His Trp Tyr Gln Gln Lys Ser His Glu Ser Pro Arg Leu
          20             25             30
Leu Ile Lys Tyr Ala Ser Gln Ser Ile Ser Gly Ile Pro Ser Arg Phe
          35             40             45
Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Ser Val
          50             55             60
Glu Thr Glu Asp Phe Gly Met Tyr Phe Cys Gln Gln Ser Asn Ser Trp
 65             70             75             80
Pro Tyr Thr Phe Gly Ala Gly Thr Lys Leu Glu Ile Lys Gln
          85             90
```

<210> SEQ ID NO 23  
 <211> LENGTH: 148  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 23

```
Met Glu Trp Ser Gly Val Phe Ile Phe Leu Leu Ser Gly Thr Ala Gly
 1             5             10             15
Val Leu Ser Glu Val Gln Leu His Gln Phe Gly Ala Glu Leu Val Lys
          20             25             30
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Pro Gly Ala Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Thr Phe  
           35                  40                  45  
 Thr Asp Tyr Asn Met Asp Trp Val Lys Gln Ser His Gly Lys Ser Leu  
           50                  55                  60  
 Glu Trp Ile Gly Asp Ile Asn Pro Asn Tyr Asp Ser Thr Ser Tyr Asn  
           65                  70                  75                  80  
 Gln Lys Phe Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser  
                   85                  90                  95  
 Thr Ala Tyr Met Glu Leu Arg Ser Leu Thr Ser Glu Asp Thr Ala Val  
                   100                  105                  110  
 Tyr Tyr Cys Ala Arg Ser Arg Ser Tyr Asp Tyr Glu Gly Phe Ala Tyr  
           115                  120                  125  
 Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ala Ala Lys Thr Thr Pro  
           130                  135                  140  
 Pro Ser Val Tyr  
 145

<210> SEQ ID NO 24  
 <211> LENGTH: 105  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 24

Gly Leu Phe Cys Ser Val Glu Arg Cys His Tyr Gln Leu Gln Ser Ser  
 1                  5                  10                  15  
 Gln Asn Leu Leu Ser Ile Val Asn Arg Tyr His Tyr Met Ser Gly Asn  
           20                  25                  30  
 Pro Pro Lys Leu Leu Val Tyr Pro Ala Leu Leu Ile Tyr Glu Ala Ser  
           35                  40                  45  
 Ile Thr Lys Ser Cys Val Pro Asp Arg Phe Thr Arg Ser Gly Ser Gly  
           50                  55                  60  
 Thr Asn Phe Thr Leu Thr Ile Asn Phe Val His Ala Asp Asp Leu Ile  
           65                  70                  75                  80  
 Phe Tyr Tyr Cys Gln His Asn Arg Gly Ser Phe Leu Pro Ser Ser Ser  
                   85                  90                  95  
 Val Gln Val Pro Arg Arg Arg Ser Asn  
                   100                  105

<210> SEQ ID NO 25  
 <211> LENGTH: 100  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 25

Asp Ile Leu Gln Ala Ser Gly Tyr Ser Phe Thr Gly Tyr Thr Met Asn  
 1                  5                  10                  15  
 Trp Val Lys Gln Ser His Gly Lys Asn Leu Glu Trp Ile Gly Leu Ile  
           20                  25                  30  
 Asn Pro Tyr Asn Gly Gly Thr Ser Tyr Asn Gln Lys Phe Lys Gly Lys  
           35                  40                  45  
 Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr Met Glu Leu  
           50                  55                  60  
 Leu Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Trp  
           65                  70                  75                  80  
 Gly Val Trp Ser Ala Met Asp Tyr Trp Gly Gln Gly Thr Thr Val Thr  
                   85                  90                  95

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Val Ser Ser Lys  
100

<210> SEQ ID NO 26  
<211> LENGTH: 90  
<212> TYPE: PRT  
<213> ORGANISM: Mus musculus

<400> SEQUENCE: 26

Asp Arg Val Ser Ile Thr Cys Lys Ala Ser Gln Asn Val Arg Thr Ala  
1 5 10 15  
Val Ala Trp Tyr Gln Gln Lys Pro Arg Gln Ser Pro Lys Ala Leu Ile  
20 25 30  
Tyr Leu Ala Ser Asn Arg Asp Thr Gly Leu Pro Asp Arg Phe Pro Gly  
35 40 45  
Arg Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile Thr Asn Val Gln Ser  
50 55 60  
Glu Asp Leu Glu Asp Tyr Phe Cys Leu Gln His Cys Asn Tyr Pro Asn  
65 70 75 80  
Glu Phe Arg Gly Cys Thr Lys Val Pro Ile  
85 90

<210> SEQ ID NO 27  
<211> LENGTH: 116  
<212> TYPE: PRT  
<213> ORGANISM: Mus musculus

<400> SEQUENCE: 27

Leu Gln Glu Ser Gly Ala Glu Leu Ala Arg Pro Gly Ala Ser Val Lys  
1 5 10 15  
Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Tyr Trp Met Gln  
20 25 30  
Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile Gly Ala Ile  
35 40 45  
Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr Gln Lys Phe Lys Gly Lys  
50 55 60  
Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala Tyr Met Gln Leu  
65 70 75 80  
Ser Ser Leu Ala Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Gly  
85 90 95  
Glu Tyr Gly Asn Tyr Phe Ala Tyr Trp Gly Gln Gly Thr Thr Val Thr  
100 105 110  
Val Ser Ser Asn  
115

<210> SEQ ID NO 28  
<211> LENGTH: 100  
<212> TYPE: PRT  
<213> ORGANISM: Mus musculus

<400> SEQUENCE: 28

Thr Ser Asp Ala Ser Leu Gly Glu Arg Val Thr Ile Thr Cys Lys Ala  
1 5 10 15  
Ser Gln Asp Ile Asn Ser Tyr Leu Ser Trp Phe Gln Gln Lys Pro Gly  
20 25 30  
Lys Ser Pro Lys Thr Leu Ile Tyr Arg Ala Asn Arg Leu Val Asp Gly  
35 40 45

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Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Gln Asp Tyr Ser Leu  
50 55 60

Thr Ile Ser Ser Leu Glu Tyr Glu Asp Met Gly Ile Tyr Tyr Cys Leu  
65 70 75 80

Gln Tyr Asp Glu Phe Pro Leu Thr Phe Gly Gly Gly Thr Lys Leu Glu  
85 90 95

Ile Lys Gln Lys  
100

<210> SEQ ID NO 29  
 <211> LENGTH: 108  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 29

Ala Trp Leu Ser Gln Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys  
1 5 10 15

Asp Thr Tyr Met His Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu  
20 25 30

Trp Ile Gly Arg Ile Asp Pro Ala Asn Gly Asn Thr Lys Tyr Asp Pro  
35 40 45

Lys Phe Gln Gly Lys Ala Thr Ile Thr Ala Asp Thr Ser Ser Asn Thr  
50 55 60

Ala Tyr Leu Gln Leu Ser Ser Leu Thr Ser Glu Asp Thr Ala Val Tyr  
65 70 75 80

Tyr Cys Ala Arg Pro Ile His Tyr Tyr Tyr Gly Ser Ser Leu Ala Tyr  
85 90 95

Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Lys  
100 105

<210> SEQ ID NO 30  
 <211> LENGTH: 104  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 30

Glu Phe His Ala Val Ser Leu Gly Gln Arg Ala Thr Ile Ser Cys Arg  
1 5 10 15

Ala Ser Glu Ser Val Asp Ser Tyr Gly Asn Ser Phe Met His Trp Tyr  
20 25 30

Gln Gln Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr Arg Ala Ser  
35 40 45

Asn Leu Glu Ser Gly Ile Pro Ala Arg Phe Ser Gly Ser Gly Ser Arg  
50 55 60

Thr Asp Phe Thr Leu Thr Ile Asn Pro Val Glu Ala Asp Asp Val Ala  
65 70 75 80

Thr Tyr Tyr Cys Gln Gln Ser Asn Glu Asp Pro Gly Arg Ser Glu Val  
85 90 95

Val Pro Ser Trp Arg Ser Asn Lys  
100

<210> SEQ ID NO 31  
 <211> LENGTH: 109  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 31

Pro Arg Ala Ser Leu Gly Val Ser Glu Thr Leu Leu Cys Thr Ser Gly

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1	5	10	15
Phe Thr Phe Thr Asp Tyr Tyr Met Ser Trp Val Arg Gln Pro Pro Gly	20	25	30
Lys Ala Leu Glu Trp Leu Gly Phe Ile Arg Asn Lys Ala Asn Gly Tyr	35	40	45
Thr Thr Glu Tyr Ser Ala Ser Val Lys Gly Arg Phe Thr Ile Ser Arg	50	55	60
Asp Asn Ser Gln Ser Ile Leu Tyr Leu Gln Met Asn Thr Leu Arg Ala	65	70	75
Glu Asp Ser Ala Thr Tyr Tyr Cys Ala Arg Ala Asn Trp Ala Phe Asp	85	90	95
Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Lys	100	105	

<210> SEQ ID NO 32  
 <211> LENGTH: 94  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 32

Ser Gly Asp Arg Val Ser Leu Ser Cys Arg Ala Ser Gln Ser Ile Ser	1	5	10	15
Asn Tyr Leu His Trp Tyr Gln Gln Lys Ser His Glu Ser Pro Arg Leu	20	25	30	
Leu Ile Lys Tyr Ala Ser Gln Ser Ile Ser Gly Ile Pro Ser Arg Phe	35	40	45	
Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Ser Val	50	55	60	
Glu Thr Glu Asp Phe Gly Met Tyr Phe Cys Gln Gln Ser Asn Ser Trp	65	70	75	80
Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Gln	85	90		

<210> SEQ ID NO 33  
 <211> LENGTH: 111  
 <212> TYPE: PRT  
 <213> ORGANISM: Mus musculus

<400> SEQUENCE: 33

Pro Ala Cys Leu Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala Thr Ser	1	5	10	15
Gly Phe Thr Phe Thr Asp Tyr Tyr Met Ser Trp Val Arg Gln Pro Pro	20	25	30	
Gly Lys Ala Leu Glu Trp Leu Gly Phe Ile Arg Asn Lys Ala Asn Gly	35	40	45	
Tyr Thr Thr Glu Tyr Ser Ala Ser Val Lys Gly Arg Phe Thr Ile Ser	50	55	60	
Arg Asp Asn Ser Gln Ser Ile Leu Tyr Leu Gln Met Asn Thr Leu Arg	65	70	75	80
Ala Glu Asp Ser Ala Thr Tyr Tyr Cys Ala Arg Ala Pro Leu Leu Tyr	85	90	95	
Tyr Ala Met Asp Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser	100	105	110	

<210> SEQ ID NO 34  
 <211> LENGTH: 102  
 <212> TYPE: PRT

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<213> ORGANISM: Mus musculus

<400> SEQUENCE: 34

Arg Leu Pro Phe Tyr Ser Leu Glu Gln Arg Ala Thr Ile Ser Tyr Arg  
 1 5 10 15

Ala Ser Lys Asn Val Ser Thr Ser Gly Tyr Ser Tyr Met His Trp Asn  
 20 25 30

Gln Gln Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr Leu Val Ser  
 35 40 45

Asn Leu Glu Ser Gly Val Pro Ala Arg Phe Ser Gly Ser Gly Ser Gly  
 50 55 60

Thr Asp Phe Thr Leu Asn Ile His Pro Val Glu Glu Glu Asp Ala Ala  
 65 70 75 80

Thr Tyr Tyr Cys Gln His Ile Arg Glu Leu Thr Arg Ser Glu Leu Val  
 85 90 95

Pro Ser Trp Lys Ser Asn  
 100

<210> SEQ ID NO 35

<211> LENGTH: 101

<212> TYPE: PRT

<213> ORGANISM: Mus musculus

<400> SEQUENCE: 35

Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Tyr Trp Met His  
 1 5 10 15

Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile Gly Met Ile  
 20 25 30

Asp Pro Ser Asn Ser Glu Thr Arg Leu Asn Gln Lys Phe Lys Asp Lys  
 35 40 45

Ala Thr Leu Asn Val Asp Lys Ser Ser Asn Thr Ala Tyr Met Gln Leu  
 50 55 60

Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Gly  
 65 70 75 80

Leu Arg His Tyr Trp Tyr Phe Asp Val Trp Gly Gln Gly Thr Thr Val  
 85 90 95

Thr Val Ser Ser Lys  
 100

<210> SEQ ID NO 36

<211> LENGTH: 99

<212> TYPE: PRT

<213> ORGANISM: Mus musculus

<400> SEQUENCE: 36

Thr Ile Leu Trp Arg Glu Gly Pro Phe Ser Tyr Arg Ala Ser Lys Ser  
 1 5 10 15

Val Ser Thr Ser Gly Tyr Ser Tyr Met His Trp Asn Gln Gln Lys Pro  
 20 25 30

Gly Gln Pro Pro Arg Leu Leu Ile Tyr Leu Val Ser Asn Leu Glu Ser  
 35 40 45

Gly Val Pro Ala Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr  
 50 55 60

Leu Asn Ile His Pro Val Glu Glu Glu Asp Ala Ala Thr Tyr Tyr Cys  
 65 70 75 80

Gln His Ile Arg Glu Leu Thr Arg Ser Glu Glu Val Pro Ser Trp Arg  
 85 90 95

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Ser Asn Lys

&lt;210&gt; SEQ ID NO 37

&lt;211&gt; LENGTH: 330

&lt;212&gt; TYPE: PRT

&lt;213&gt; ORGANISM: Homo sapiens

&lt;400&gt; SEQUENCE: 37

Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys  
 1 5 10 15

Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr  
 20 25 30

Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser  
 35 40 45

Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser  
 50 55 60

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr  
 65 70 75 80

Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys  
 85 90 95

Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys  
 100 105 110

Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro  
 115 120 125

Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys  
 130 135 140

Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp  
 145 150 155 160

Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu  
 165 170 175

Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu  
 180 185 190

His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn  
 195 200 205

Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly  
 210 215 220

Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu  
 225 230 235 240

Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr  
 245 250 255

Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn  
 260 265 270

Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe  
 275 280 285

Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn  
 290 295 300

Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr  
 305 310 315 320

Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 325 330

&lt;210&gt; SEQ ID NO 38

&lt;211&gt; LENGTH: 106

&lt;212&gt; TYPE: PRT

&lt;213&gt; ORGANISM: Homo sapiens



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&lt;400&gt; SEQUENCE: 38

Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln  
 1 5 10 15  
 Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr  
 20 25 30  
 Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser  
 35 40 45  
 Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr  
 50 55 60  
 Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys  
 65 70 75 80  
 His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro  
 85 90 95  
 Val Thr Lys Ser Phe Asn Arg Gly Glu Cys  
 100 105

The invention claimed is:

1. An antibody or a fragment thereof which has immunological reactivity with a CAPRIN-1 protein, the antibody or the fragment thereof comprising a heavy chain variable region comprising complementarity determining regions of SEQ ID NOs: 5, 6, and 7 and a light chain variable region comprising complementarity determining regions of SEQ ID NOs: 9, 10, and 11.

2. The antibody or the fragment thereof according to claim 1, wherein the antibody is a human antibody, a humanized antibody, a chimeric antibody, a single-chain antibody, or a multispecific antibody.

3. The antibody or the fragment thereof according to claim 1, wherein the antibody or the fragment thereof is conjugated with an antitumor agent.

4. A pharmaceutical composition for treatment of CAPRIN-1- cancer, comprising an antibody or a fragment thereof according to claim 1 as an active ingredient.

5. The pharmaceutical composition according to claim 4, wherein the cancer is breast cancer, kidney cancer, pancreatic cancer, large bowel cancer, lung cancer, brain tumor, gastric cancer, uterine cervix cancer, ovary cancer, prostate cancer, urinary bladder cancer, esophageal cancer, leukemia, lymphoma, fibrosarcoma, mastocytoma, or melanoma.

6. A pharmaceutical combination for treatment of CAPRIN-1- expressing cancer, comprising a pharmaceutical composition according to claim 4 and a pharmaceutical composition comprising an antitumor agent.

7. A DNA encoding an antibody or a fragment thereof according to claim 1.

8. A method for treating CAPRIN-1-expressing cancer, comprising administering an antibody or a fragment thereof according to claim 1.

9. The antibody or the fragment thereof according to claim 2, wherein the antibody or the fragment thereof is conjugated with an antitumor agent.

10. A pharmaceutical composition for treatment of CAPRIN-1-expressing cancer, comprising an antibody or a fragment thereof according to claim 2 as an active ingredient.

11. A pharmaceutical composition for treatment of CAPRIN-1-expressing cancer, comprising an antibody or a fragment thereof according to claim 3 as an active ingredient.

12. A pharmaceutical combination for treatment of CAPRIN-1-expressing cancer, comprising a pharmaceutical composition according to claim 5 and a pharmaceutical composition comprising an antitumor agent.

13. A DNA encoding an antibody or a fragment thereof according to claim 2.

14. A method for treating CAPRIN-1-expressing cancer, comprising administering an antibody or a fragment thereof according to claim 4.

15. A method for treating CAPRIN-1-expressing cancer, comprising administering an antibody or a fragment thereof according to claim 6.

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